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	1 - 490	1 - 464	1 - 480	1-418	1 - 188	1 - 158	1 - 113	1 - 276	1 - 187	1 - 321	1 - 216	1 - 267	1 - 158
	968455	927811	950732	868109	529572	529571	529573	861074	529574	529577	529578	968440	529579
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1 - 350	1 - 255	1 - 66	1 - 288
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-	8091	956253	1 - 360	15 - 374	
HMWIM05 8	8092	861094	1 - 567	15 - 581	AA160427, AI912401, AA228442, AW272815, AA972523, AA484892, AA728911, AA449923,
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	15 - 722	15 - 625	15 - 332
	1 - 708	1 - 611	1 - 318
	959512	964701	767332
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	15 - 368	15 - 487	15 - 355	15 - 161	15 - 599
	1 - 354	1 - 473	1 - 341	1 - 147	1 - 585
	861091	663972	657260	913689	419821
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IIMWBH29	8145	927321	1 - 625	15 - 639	T74339, AA007592, W90626, AA680304, AA954254, AI133697, W60511, AA007389,
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HMVDV55	8150	811616	1 - 397	15-411	R56378, H05003, and R83285.
НМУДО02	8151	919057	1 - 342	15 - 356	AA663501, AA862536, AW265642, and AW275747.
HMVDL45	8152	717384	1 - 827	15 - 841	H09382, R17471, Z44450, and F07290.
HMVDJ60	8153	877124	1 - 774	15 - 788	N46872, AA046610, AI589041, AI638264, AI888302, AW206234, AW135977, AI580058,

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	1 - 584	1 - 440	1-870	1 - 506	1 - 435	1 - 372	1 - 695										1 - 432	1 - 501		1 - 364	1 - 439	1 - 797	1 - 406	1 - 788				٠						
	661692	786118	780366	790025	726546	716410	888180	•					_	_			726515	732985		761400	861210	751600	861245	926800										
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	HMVBV16	HMVBU89	HMVBO83	IIMVBJ91	HMVBF52	HMVBF44	HMVAT54	•									HMVAT52	HMVAP56		HMVAF71	HMVAF52	HMVAC67	HMSPH53	HMSPE04								_		

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	8173	8175	8176	8177	8178	8179
	HMSPA30	HMSOJ88	HMSOJ01	нмѕоноз	HMSNP06	HMSNG04

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15 - 432	15-131	15 - 724										-1.																							
1 - 418	1-117	1 - 710																																	
963431	861336	861338	•	-														•																	
8180	8181	8182																																	
HMSMY10	HMSMW91	HMSMW64																																	

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	15 - 609	15 - 649	15 - 503	15 - 182	15 - 527	15 - 576	15 - 498	15 - 2169	15 - 560	15 - 394
	1 - 595	1 - 635	1 - 489	1 - 168	1 - 513	1 - 562	1 - 484	1 - 2155	1 - 546	1 - 380
	861341	927097	861345	953390	861372	495770	783032	852496	775416	954003
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	HMSMO59	HMSMN71	HMSMN26	HMSLH07	HMSKQ08	HMSKI124	HMSJX84	HMSJW48	HMSJU80	HMSJQ40

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		15 - 270	15 - 64	15 - 1093	15 - 434	15 - 583	15 - 475	15 - 468	15 - 647						-
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	1 - 815	1 - 365	1 - 506	1 - 326	1 - 339	1 - 390	1 - 339	1 - 1310	1 - 324	1 - 499	1 - 265	1 - 503	1-357	1 - 723	1 - 1336	1 - 383
	465835	765600	699241	686663	669671	920079	615218	935151	967007	753038	495778	890296	717808	709471	741834	959461
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HIMISHEI!	6779	607700	1 - 833	608 - CI	1/9224, 1/9/33, A18/8808, A18/8839, AW160301, W2/440, D59610, AW361175, T03269,
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					D59787, AW178893, D80366, D50979, D50995, D80164, D80024, D80378, D80045, D51250,
					C14331, C15076, C14389, C14014, AW378532, C75259, D51060, AW177440, AA305409,
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	15-696	15 - 324	15-379	15 - 569	15 - 587	15 - 430	15 - 839
	1 - 682	1 - 310	1 - 365	1 - 555	1 - 573	1 - 416	1 - 825
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	15 - 462	15 - 184	15 - 422	15 - 1988
	1 - 448	1 - 170	1 - 408	1 - 1974
	745230	888326	959773	068616
	8228	8229	8230	8231
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2 2 2 2 3 3 4 4 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5 5	15 - 308 AI	15 - 151	15 - 223	15 - 544 T8	15-366 H60137	15 - 200 AJ	15 - 235	15 - 276 H	15 - 431 Az	15 - 166 N56866	15 - 329	15 - 353 H: A) Al	15 - 366 A. T.3	A
	1 - 294	1 - 137	1 - 209	1 - 530	1 - 352	1 - 186	1 - 221	1 - 262	1 - 417	1 - 152	1-315	1 - 339	1 - 352	_
	959774	753635	384695	960147	706366	842040	751865	660654	266606	968435	559991	574950	927804	
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·	HMSFN08	HMSEN68	HMSEN13	HMSEN08	HMSEM34	HMSEL64	HMSEG67	HMSDQ15	HMSDL70	HMSDK10	HMSDI34	HMSDG12	HMSDG04	

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8255 752145 1 - 464 15 - 585 8256 880721 1 - 571 15 - 585

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15-771	15-377	15 - 141	15 - 190	15-85	15 - 462	15 - 1325			<u>-</u>																								
1 - 757	1 - 363	1 - 127	1 - 176	1-71	1 - 448	1-1311																											
628709	796724	792970	780229	733772	784576	958802																											
8257	8258	8259	8260	8261	8262	8263																					_						
HMSBU24	HMSB095	HMSBM92	HMSBI82	HMSB156	HMSBH85	HMSBH08														•													

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	15 - 462	15-919	15 - 566		
	1 - 448	1 - 905	1-552	1 - 364 1 - 405 1 - 393	
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	8264	8265	8266	8267 8268 8269	
	HMSBH07	HMSBF92	HMSBC31	HMSBA43 HMSAV84 HMSAV65	

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	15 - 637	15 - 395	15 - 360	15 - 225	15 - 188	15 - 379	15 - 464	15-358	15 - 256	15 - 386
	1 - 623	1 - 381	1 - 346	1-211	1 - 174	1 - 365	1 - 450	1 - 344	1 - 242	1 - 372
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	8270	8271	8272	8273	8274	8275	8276	8277	8278	8279
	HMSAT76	HMSAQ95	HMSAQ58	HMSAP36	HMSA042	HMSAH80	90ЛООМН	HMQDQ42	HMQDL40	HMOC736

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15 - 473	15 - 445		15 - 398		15 - 149	15 - 194	15 - 433	15 - 279	15-319										15-473	000	15 - 3/0	15 - 304	15 - 345	15 - 745	15 - 353	15 - 242	15 - 482	15 - 343	15-327	15 - 600	15 - 452
1 - 459	1 - 431	1 - 486	1 - 384	1-123	1 - 135	1 - 180	1 - 419	1 - 265	1 - 305				-	-					1 - 459	,	1 - 350	1 - 290	1 - 331	1 - 731	1 - 339	1 - 228	1 - 468	1 - 329	1 - 313	1 - 586	1 - 438
574068	712966	974683	385544	719201	711586	932363	754188	728851	963370										779856	00000	921233	682099	709222	781863	600400	855919	796276	913715	754902	625366	875272
8280	8281	8282	8283	8284	8285	8286	8287	8288	8289										8290	1000	1678	8292	8293	8294	8295	-	8297	8538	8299	8300	8301
HMQCZ18	HMQCX42	НМQСV29	HMQCN17	HMQCM46	HMQCM41	нмосмоя	НМОСП69	HMQCH54	н.мосвоз										HMQB:N40	The street of the	HMF1A56	HMPTA28	HMPSC38	HMPBB84	HMOAD90	HMOAD60	HMMCL96	HMMCI01	HMMCH69	HMMCH09	HMMCD83

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HMMBV08	8308	958987	1 - 450	15 - 464	AI939575, and AI362923.
HMMBU13	8309	597147	1 - 444	15 - 458	H65100, AC004686, AF201333, and AF081568.
HMMBT64	8310	746399	1 - 408	15 - 422	T96528, and AA543014.
IIMMBT30	8311	841959	1 - 607	15 - 621	AA280370, R58990, II71497, AC005050, and AL031985.
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	HMMBQ08	HMMBO49	HMMB028	HMMBN76	HMMBN71	HMMBN54	HMMBM8	HMMBL55	HMMBH64	нммвғ86	HMMRF72

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					AA218811, AA297863, AA296837, R95168, N86960, R39644, T10507, AA485829, AI342703,
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HMCHS34	8351	703691	1 - 483	15-497	T86231, T83384, and T86130.
HMCHK58	8352	752431	1 - 574	15 - 588	AA160383, AI818161, AI962470, AI655470, AI201401, AI962522, AI809442, AA757199,
					AA954406, AI557476, W46315, AI263049, AI557453, AA332432, AI867723, AA383667,
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HMCGV66	8354	787512	1 - 203	15-217	A1159933, A1971968, AA906691, A1183889, W79831, N92399, AA557293, AI350318, AI017848,
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					AA937/86, AA988459, AA404424, A1081962, W19157, A1129889, AA935718, A1066610, and AF111170.
HMCGF39	8355	856222	1 - 484	15 - 498	W27128, and H14722.
HMCFR57	8356	734404	-1 - 405	15-419	AI304652, AI347073, AI343275, AI261508, F26025, AA936207, N66083, AA769680, AI867524,
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					AI025107, AA806009, AA007654, AI033216, AI399750, T06491, AA917469, AA134955,
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					AA309446, AA424656, AA492384, AF132945, and AF112204.
HMCFM04	8358	927002	1 - 529	15 - 543	AA569402, AW070532, N38778, AW190781, AW340905, AA535216, AW023111, AA773463, AW0337905 AA714110, AA828637, AA524616, AI674840, A1755202, A1066646, AW77224
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AA225406, AA456924, AI061313, AA632765, AW277174, AI635028, R92658, AI755214.
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AA558404, AW081485, AA548610, AI355246, AW130427, AI583466, AA829065, AW075132, H71678 A 1945605 A 1946605 A 19466
AW274191, AA557982, AA581525, AI801649, AI301373, AA536040, AI446452, AW167330,
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A A O A A A O A A A O A A A O A A A O A A A O A A A O A A A O A A A O A A A A O A A A A O A	15 - 354 AI8 AI3	15 - 374 AJ3	15 - 565 AL	344	15 - 409 W21990	- 546	- 647	15 - 545 AI3	15-460 H87	15-541 AA	AA	Z30	AIS	R29	YT(AL.	- AL	AIS	AL	- AT	AL	AL	AL	AL	<u>Ā</u>	AL	- AT	AE	AL	AL
	1 - 340	1 - 360	1 - 551					1 - 531	1 - 446	1 - 527				•	-	•					••					***				
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	HMCDN93	HMCDC70	HMCAR04	HMAL277	HMALK06	HMAJZ39	HMAJL09	HMAIU90	HMAIQ20	HMAIH15																				

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HMADF74	8377	944465	1-1131		AA311043, H11578, R59755, AA927842, AA199784, AI000363, AI126309, R13187, AI693684, and AI 096806.
HMACZ84	8378	975349	1 - 340	15 - 354	
HMACV54	8379	922354	1 - 409	15 - 423	AI000925, AI261943, AA614341, AA532388, AI000930, AI742836, AW376738, AW136154, AW376570, and E08096.
HMABT55	8380	753560	1 - 444	15 - 458	N50809, AA931135, AA972697, AA401345, AI024270, AA873221, AW082806, AW025001, AI752742, AI38745, AI277497, AW452978, AI554935, H50235, AI911227, AI700463, AW303800, AI275531, AI241950, AI760168, AI095901, AA618048, AI936652, R09749, AA936697, AI199598, AI206725, AI830389, AA084391, AA987748, H64022, H06700, AI926722, AA988399, AA838228, AI357218, AI362511, AI362439, AA937768, AI879499, AI693589, AI366339, AI36539, AI69389, AA663409, H20195, AW083589, AI963935, AA843550, H64785, AI243967, AA725384, AW449564, AW450568, AW339993, W96140, W92486, H41770, and AA018786.
HMABD66	8381	388074	1 - 370	15 - 384	
HLYHN64	8382	970747	1-577	15 - 591	N53558, AI078352, H49897, W00521, N30989, N57223, AA314759, AA431398, AA972343,
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HLYHL55	8384	731447	1 - 447	15 - 461	R32479.
нгунк64	8385	746310	1 - 456	15 - 470	H53572, AA745885, H28445, AI494060, AI676167, AA889745, AI866665, AI245303, AA703022, AI016995, AW264349, AI254741, AI287476, AW160916, AA580663, AI345688, AA742390, AA641818, AL138406, AI590043, AI309380, AI567846, AI537244, AW075382, AI538885, AW151136, AI432644, AI500061, AW079432, AW022808, AI859991, AI953765, AL037454, AI623941, AW022636, AI287233, AI702527, AA766077, AW161579, AI540674, H89138, AI587121, AW304497, AW268759, N29277, AI673278, AI918449, AL118781, AI473451, AA811142, AI538764, AI868204, AI866461, N63128, AW161156, AL037602, AI446373, A1969655, AW020095, AI499890, AW021717, AI890907, AI345745, AA676361, AI929108, AW020419, AL039086, AL047344, AI581033, AI961278, AI537274, AI432570, AI686576, AW020693, AI340603, AL037682, AI537677, AI621341, AI524654, AI888621, AI473536, AL135024, AI284517, AI560545, AI923989, AI241901, AI783530, AW020397, AW192037, AW020337, AW020337, AW020337, AW020337, AW020337, AW020337, AW020337, AW020337, AW020332, AI871251, AW16388, AI472484, AI633125, AI872423, AI636619, AI533645, AI636631, AI915291, AW152182, AW274318, AI582932, AI872423, AI540606, AI839189, N52016, AI533800, AW195253, AI309306, AW104141,

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					AA922534, AA533814, W16698, AW169172, AA654163, AA412735, N69522, AI440188,
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HLYEA70	8431	943886	1 - 739	15 - 753	W79930, AA233418, T87364, AI765694, AI042017, AI672297, T87365, R94396, AI082728,
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935611 1-527 15-541 AA736563, AA642445, AA838199, AA724333, AA669153, AA380354, AI537772, AI114701, AW4935563, AW493545, AW319180, AJ950671, AW151713, AW32099, AA665146, AW022161, AW304580, AW490545, AW301385, AA704393, AA832077, L35667, AC006538, AP000557, AL009031, AF135187, AC004020, AA832077, L35667, AC006739, AP000557, AL009031, AF135187, AC004020, AC0062185, AC007199, 283987, AP000605, AL050348, AL03174, AC004020, AC0062185, AC00236, AC007199, 283987, AP000605, AC006196, AC006196, AC006196, AC006197, AC005211, AF207550, AC006036, AC006312.79, AC005191, AC002211, AF207550, AC006036, AC006312.79, AC005191, AC006211, AF207550, AC006246, AC005519, AC005211, AF207550, AC006246, AC005519, AC005211, AF207550, AC006246, AC005519, AC005211, AF207550, AC006246, AC005519, AC005211, AC006291, AC006911, AC006291, AC006911, AC005291, AC006911, AC006911	8473 767225 1 - 342 15 - 356 AC004242.	8474 699337 1-188 15-202 A1926931, T92757, and AC004984.	/82.555 1 - 5/0 15 - 564 AA095492, AW404010, W15542, AA061452, 16/155, AA500169, A1504007, 16/1262, A1504007, A15	8476 856469 1 - 430 15 - 444 AA703693.	8477 678179 1-470 15-484 AA193337.	 	757235 1	754229 1 - 329 15 - 343	815950 1-365 15-379 AL10962	854034 1 - 330 15 - 344	8483 964834 1 - 651 15 - 665 AA487424, AA578890, and AI524037.
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HLYBA06	HLYAX75	HLYAX32	HLYAUIS	HLYAS35	HLYAR25	HLYAQ92	HLYAQ70	HLYAQ69	HLYAN68	HLYAN21	HI YAN10

AA847181, AA470533, AA484373, AW243954, AI821918, AI821881, AI268334, AI268336, AA805363, AA507912, AI470431, AA582218, AA715805, AA582229, AA600869, AA605272, AL038936, AI605038, AI805547, AA6682238, AA527209, AI470437, AI500453, AI500852, AA605272, AL038936, AI805236, AA668022, AA501718, AI922599, AW277171, AZ50083, To6681, AW702259, AW102846, AA468022, AA311208, AL0312599, AA47775, AI801482, AA731235, AW702359, AA731823, AA7422998, AI817976, AA407391, AI669046, N84446, AA651632, AA702391, AA730823, AA74230998, AI817976, AA605231, U07561, AC005003, AC007666, AL031005, AC005387, AC004821, AL031232, AC005231, AC005314, AC00589, AC005015, AL03166, AC005919, AL133245, AC007225, AC005011, AF165926, AC007686, AC004685, AC005911, AC00599, AL133448, M63796, AC004480, AC004376, AC005384, AC005364, AC004880, AC004488, AC004883, AC004488, AC004488, AC004883, AC004488, AC004883, AC004883, AC004883, AC004883, AC004883, AC004883, AC004914, AL0312874, AL031688, AL031377, AC005880, AP000117, AL022238, AC0047216, AL031689, AL031387, AL031883, AL031283, AL031283, AL031283, AL031283, AL031283, AL031283, AL031283, AC004381, AC00341, AL032234, AL030263, AC004381, AC00359, AC003491, AL032234, AL03066, AC004383, AC00359, AC003484, AC00359,	AA809368, AW190881, and Al240461.	T92768, and AA682689.	AA013385, W96162, H84095, AA059069, AA054036, H84089, H86124, AA018232, AA018236,	W24486, C06267, R42858, AA978262, AA505146, AI636820, AI143214, AA045249, AI677860, AA491378, AA255801, and AL031297.	
15 - 457	15-341	15-370	15 - 444	.15 - 436	15 - 441
1 - 443	1 - 327	1 - 356	1 - 430	1 - 422	1 - 427
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	8486	8487	8488	8489	8490
HLYAL22	HLYAL11	HLYAK36	HLYAK34	HLYAJ49	HLYAG59

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15 - 803	15 - 360	15 - 639	15 - 460	15 - 877	15 - 504					_					•										_											
1 - 789	1 - 346	1 - 625	1 - 446	1 - 863	1 - 490																															
708613	856560	708620	780903	708027	662409																															
8491	8492	8493	8494	8495	8496																															
HLTHS37	HLTHS02	HLTHC37	HLTGT83	HLTGT36	HLTGS17															-																

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IDO		HLTGR28 8497

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HLTGP47	8498	719975	1 - 433	15 - 447	A64136, and A68321. AW393850 AA194696, and AW274557
HLTG089	8499	786674	1 - 594	15 - 608	R24848, and Z43303.
HLTGN92	8500	791185	1 - 358	15 - 372	R83296, H61528, R06428, and AA497003.
HLTGA40	8501	710728	1 - 401	15 - 415	H12953, AA436993, and AA426378.
HLTEZ30	8502	960911	1 - 338	15 - 352	T85564, T84409, AA361578, and R08443.
HLTEZ19	8503	671610	1 - 705	15 - 719	N98660, AI052555, AI033178, R16888, and AA437100.
HLTEX27	8504	927578	1 - 336	15 - 350	N95142.
HLTEV66	8505	772965	1 - 506	15 - 520	AA046570, AI912976, AI810637, AI400892, AI420896, AA317499, N44648, AI085349, T49602,
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HLTEP45	8506	723375	1 - 524	15 - 538	AA159675, and AC007068.
HLTEM25	8507	678685	1 - 508	15 - 522	R14032, AA825482, H95342, H70614, and AC004921.
HLTEGS1	8208	494330	1 - 605	15-619	
HLTEG37	8509	417243	1 - 363	15-377	W40567, AL040311, AA905330, AA677615, AW205363, and D80009.
HLTEC83	8510	574630	1-213	15-227	
HLTEC35	8511	574632	1 - 266	15 - 280	
HLTEA20	8512	006699	1 - 582	15 - 596	AA007473, W93011, Z44985, AA430251, and N71668.
HLTDR43	8513	715529	1 - 664	15-678	A1953354, H16401, H00613, F09067, R82764, Z38624, and T17430.
HLTD117	8514	577379	1 - 657	15 - 671	W56369, I184793, AA913919, AA485021, AC007387, AC006543, and AC006544.
96НСІЛН	8515	796560	1 - 275	15 - 289	N72412, and AI689177.
HLTDG59	8516	739699	1 - 340	15 - 354	T81367, AA372287, R50286, W30782, and AL137669.
HLTDG30	8517	693329	1 - 345	15 - 359	H85024.
HLTDB38	8518	456860	1-418	15 - 432	R95058, and R83706.
HLTCZ55	8519	732385	1-312	15 - 326	AA429774.
HLTCX39	8520	705648	1 - 432	15 - 446	AA157834, and AF095703.
HLTCT77	8521	772458	1 - 356	15-370	H84601, and AW379855.

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H84886.	AAS48022, and AC005740.	AA633942,	R12467, and AB020672.	A1150147, and A1362692.	R67858, and AC008033.	H54393, AA476547, C02267, and AA445910.		AI569948.	H38593, AA361336, Z99571, and Z83820.	AA521184, AA766096, AA743785, and AA361334.	R18278, R17529, AA361271, AA019429, T66081, F12334, and AW025272.	W78875, AA463793, R10150, AA814668, AA779112, R10157, AI351903, AA361232, and AC004554.	AA150350, and AA361174.	AA282169, AA987588, and AA360588.	N45611, and AA360686.	W55847, T65532, F11980, AA255754, AA360399, and N45924.	R73095, R26859, R73096, AA360390, and R27089.	AA828265, and AW262531.	A1076363, AA001640, and AA001228.		N55558, AA280889, AA095152, T90017, and AC008124.	T39190, and T40464.		W42656, and AA077151.	R22164, A1248937, A1084822, A1249135, and AA897402.	AA837202, and Z82189.	AA716681, AI361749, N77694, H65481, AA005280, H64888, AA005196, N58327, AA747865, R94521, AA677973, A1032897, R93670, II73820, H54422, and AC012627.	AL120141, AF111168, AC005081, AL109798, AC004883, AL022476, AC006441, AC005071,	AC005796, AC009263, AC002369, AL049569, AC006064, AL050341, Z98946, AC002430,	U62293, AC006312, AL109628, AC006211, AC005189, AC005736, AC007384, AC006023,	M29930, AC005057, AC004638, Z81364, AP000689, M29929, AP000248, AC005317, Z68756,	AC004990, AC005225, AC005049, Z93930, AC005562, AP000210, AC005089, AC005332,
15 - 387	15 - 349	15 - 1187	15 - 501	15 - 317	15 - 367	15 - 356	15 - 449	15 - 249	15-378	15 - 527	15 - 422	15 - 667	15-380	15 - 667	15 - 227	15 - 649	15-391	15 - 498	15 - 489	15 - 472	15 - 220	15-616	15 - 506	15-210	15 - 386	15-510	15 - 633	15-356				
1-373	1 - 335	1 - 1173	1 - 487	1 - 303	1 - 353	1 - 342	1 - 435	1 - 235	1 - 364	1 - 513	1 - 408	1 - 653	1 - 366	1 - 653	1 - 213	1 - 635	1 - 377	1 - 484	1 - 475	1 - 458	1 - 206	1 - 602	1 - 492	1 - 196	1 - 372	1 - 496	1-619	1 - 342				
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	1 - 473	1 - 432	1 - 445	1 - 448	1 - 353	1 - 171	1 - 430	1 - 744	1 - 220	1 - 439																						
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HLMFE79	8561	531213	1 - 388	15 - 402	
HLMFE17	8562	531218	1 - 137	15 - 151	
HLMFD62	8563	754076	1 - 322	15 - 336	AI797595, AW007013, AI498775, AI498624, H09241, AA935139, AA948609, R44566, and
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HLMDX01	8264	921340	1 - 401	15-415	AAZ32886, and 181241.
HLMDH08	8565	960055	1 - 814	15 - 828	A1423013, AW058211, AA700642, AW027855, AI709117, AI673385, AW339470, AI161207,
					AA694181, A1276286, A1216865, H54304, H54097, T99459, T98863, AA669188, AW386076, and AB033044
HLMDC26	8566	684911	1 - 452	15 - 466	N23041, AW380413, AA078090, AL048960, N28704, AC005088, AF030453, and AJ007798.
HLMCL59	8567	529908	1 - 586	15 - 600	AA742279, N80170, W03637, and AL031289.

N77018, A1916826, AW236861, AA429686, and AI623392.	AA51549, AA601425, AW089625, AA639170, AA620554, T47745, AA516058, AI056526, AW102811, AL040382, T06734, AI110760, AA59341, AA523833, AA463204, AW40534, AW102811, AL040382, T06734, AI110760, AA593415, AA6402727, AF190465, U52112, AC00412, AC006193, AC005284, AL113163, AC004815, AC004223, AC007216, AC00586, U95742, AC004815, AC005891, AL1031033, AC004832, AC005808, AP000557, AC005417, AC005081, AL09907, AP000045, AC005081, AC003101, AC00546, AC003101, AC003411, AC009247, AC007055, AC002507, AL099761, AC007086, AC007313, AC009511, AC009247, AC007055, AC002507, AL099761, AC007086, AC00312, AF196969, AC003511, AC004991, AC007061, AL049780, AC005707, AC003077, AC003077, AC003077, AC003077, AC003070, AL089780, AC005307, AC003077, AC003083, AC003077, AC003077, AC003083, AC003077, AC003077, AC003083, AC003083, AC003077, AC003077, AC003077, AC003083, AC003083, AC003083, AC003077, AC003084, AC003077, AC003083, AC	AC007688, AL121603, AC006071, AC005585, AL023807, AC005180, AC004149, AC006121, AC005562, AC005940, and AP000248.	H72138, and AA594731. AA505108. and N72195.	
15 - 439	15 - 860		15 - 468	
1 - 425	1 - 846		1 - 454	
741635	879359		730952 560848	
8958	8569		8570	
HLMC155	нгмвх91		HLMBQ54 HLMB048	

HLMBN04	8572	615620	1 - 340	15-354	T77272, F13388, and AF052103
HLMAV56	8573	733708	1 - 540	15 - 554	AA234263, and AA262980.
HLMAI82	8574	523694	1 - 100	15 - 114	
HLMAE86	8575	852563	1 - 543	15 - 557	AA056759, A1870884, A1184256, AA287354, AL048819, AA780049, AA658119, AW083033, AW131126, AA812201, AA523491, A1955653, AA102847, A1420177, A1658829, A1860578, AA058437, AW439401, AA700175, AA527995, AW439499, A1299102, A1471747, AA102846, A1767122, AA884538, A1859388, A1813491, AA004770, AA973515, A1016255, AA452810, H22045, A1686317, AW753434, A AA57975, H09535, A178601, A1865742, A A A A A A A A A A A A A A A A A A A
					AI916285, AA310886, AI263292, R41868, AI868055, AA809214, AW439033, AA994606, AA682833, AA004702, AA287405, and AI597574.
HLLCD85	8576	783454	1-515	15 - 529	AI478306, AI161172, AA554537, W68698, AI380170, AI085355, AI796561, AI571389, T70262, AA188734, N92493, AW385352, AI216082, AA188572, AI141460, AI357387, AA489805, AA398636, AF108460, AF108461, AC020663, AF108455, and AF108454.
HLKDC76	8577	953502	1 - 1087	15 - 1101	AA161515, AW250345, AW364752, AW248279, AA173349, AA410730, T09386, N28872, AA281341, N33896, AA312252, AA160094, T08124, Z45256, F07609, AA452741, R69813, AA380203, AA256741, N57212, Z43138, A1017337, F07095, T32695, W26853, T35905, AW204528, R14455, AI948689, AA682474, AI378863, AA378888, W27662, AI935695, AI928405, and AA910271.
HLKDB78	8278	773172	1 - 459	15 - 473	AW298414, N39216, AW004711, AW051893, AI375134, AI917477, AI652977, AI291261, AW002993, AW003527, AI016680, AI676093, AA668759, R49090, AI351891, AI498667, AI203099, AA906803, AI356617, AI950955, Z40923, AI783643, AI418750, AI189132, AF069532, and Y16521.
HLKDB02	8579	918870	1 - 458	15 - 472	T55231.
HLEAN30	8580	843294	1 - 442	15 - 456	
HLCDA67	8581	887375	1 - 619	15 - 633	
IIKBBV11	8582	965170	1 - 528	15 - 542	AA249141, AA317762, AA323439, R68885, AA657752, AA845804, AA443184, AL138096, AC006998, AC005406, AL022396, AC003029, AC004381, AP000008, AP000704, AL021367, AL031595, AC004106, AC008012, AC004087, AP000503, AF134726, AL034421, AL031274, AC002528, AC006536, AC006077, AL133246, AL034371, AL031733, X54486, AL031650, AC004033, AC007731, AC005500, AL034345, AL022098, AC004832, AL079299, M26434, AC004231, AC004383, AC005243, AF099810, AC003101, AC020663, AC005354, AC005722, and AC005261.
HKBBP05	8583	928316	1 - 539	15 - 553	A1436366, and H81947.
HKBAL06	8584	934160	1 - 527	15 - 541	A1039905, AA826224, and AA911597.
HKBAJ12	8585	969644	1 - 648	15 - 662	AA742235, AA742234, AA769592, AA828385, and AA804905.
HKBAB02	8586	918568	1 - 584	15 - 598	AA524543, H28219, AA482728, and AC006486.

HJUBA59	8587	900888	1 - 631	15-645	AI971330, A1266060, AI870369, AA025208, N48096, AI673475, AW058505, AI864572, R08870,
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HJTRB51	8888	470587	1 - 486	15 - 500	AA648882, and AA357310.
нжрк80	8589	418622	1 - 902	15-916	AA210877, AI797681, AI970134, W86482, AA070750, AA214621, AA977114, W86632, AA252120, AA902411, AA742986, AA242915, AA579700, AI468181, AA376316, AW015483, AA811628, AI928274, AI799954, AA040813, AA213789, and AF001549.
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HJPD116	8591	518105	1 - 343	15-357	H29753, AA011703, AI341972, AA477868, AI813605, AA581094, AW027687, AW193238,
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HJPDE23	8592	574115	1 - 231	15 - 245	
HJPDD82	8593	857401	1 - 990	15 - 1004	AA455589, A1091855, AA988340, AA161359, H42568, R07607, A1015867, R07600, R07602,
		-+			AA441951, A1936472, and A1291961.
HJPCW09	8594	536677	1 - 234	15 - 248	Z44259, H07973, AB015348, AL137622, AL133072, AL133048, AL133040, AJ132440, AJ243706, and AF087481.
HJPCV54	8595	733607	1 - 575	15-589	AA402378, AA464262, AI885402, AI829944, AI921923, AW008540, AI074307, AI005213,
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					AW242087, AA873160, AI362816, AA857987, AA745265, AI991184, AI222764, AI972124,
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	8296	8597	8298	8599	0098		8602	8603	8604	8605	9098	2098										
	HJPCS54	НЈРВР22	HJPBA04	HJPAZ91	HJPAY56	HJPAX11	HJPAT79	HJPAG38	HJPAA60	HJKSC56	HJBDM21	нлврг70										

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		15 - 579	15 - 554	15 - 331	15 - 227
		1 - 565	1 - 540	1-317	1 - 213
		973120	901275	780456	806332
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		HHETK38	ннетн36	HHESV83	IIHESV71

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8998	6998	8670	8671	8672	8673	8674	8675
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	1 - 2504	1 - 1057
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	8676	8677
	HHEPT05	ннертоз

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HHEPM69	8679	588235	1-372	15 - 386	

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15 - 828 A	15 - 543 A	15-477 A	15 - 367 N	15-535 T		15 - 403 A	< <	<u> </u>	72	∀	<u> </u>	<u> </u>	<	<u> </u>	<u> </u>	<u> </u>	<u>~</u>	-	¥ ·	Α .	€ ◆	2	- -	Y Y	<u> </u>	<u> </u>	<u> </u>	Ψ_	A U
1 - 814	1 - 529	1 - 463	1 - 353	1 - 521	1 - 361	1-389		*				,	-	-															
877268	925436	956241	588155	744995	919775	887661						-																	
8680	8681	8682	8683	8684	8685	9898		·	•																				·
HHEPG62	ннерд59	HHEPF21	HHEOG81	HHEOA63	HHENY02	HHENWII																							

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HHENO08	8687	959262	1-431	15 - 445	H68621, H80833, and AC004622.
HHENK21	8898	858290	1-798	15 - 812	AA631620.
HHENJ29	6898	90576	1 - 488	15 - 502	A1357399, A1813392, A1769194, AA490985, AA648531, AA971925, N90859, A1074897,
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HHEND53	8690	728111	1 - 500	15 - 514	II22946, and Z38757.
HHENC71	8691	757844	1 - 530	15 - 544	W95913, and AC003090.
HHEMZ66	8692	586737	1 - 432	15 - 446	AW340990, AW270170, AA913381, AW392670, AL119341, AL119319, AL119457, AL119324,
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HHEMW78	8693	773507	1 - 725	15 - 739	AA251113, AA766300, AA651948, AA703557, R94607, AL119319, R94247, AL042551,
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HHEMV01	8694	916281	1 - 262	15 - 276	R95819, H61040, H61032, and AC003681.
ннем068	8695	752326	1 - 140	15 - 154	AC006991, AC008175, and AC007967.
ннем048	9698	493733	1 - 534	15 - 548	
HHEMN37	8697	708468	1 - 205	15-219	
HHEMK77	8698	772115	1 - 255	15 - 269	

	789443	1 - 458	15 - 472	AC004047. R38756. AA169642, AA807905, AW021148, AA746636, W76332, AA041299, W72247, Z40683.
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	15 - 462	15 - 719	15 - 298	15-414	15-471			15 - 1827					•														
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	8703	8704	8705	8706	8707			8208								_											
	HHEMI09	HHEMG02	HHEMG01	HHEMD56	HHEMC41			HHEMC10													-···-						

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HEIAU39	8791	861503	1 - 354	15 - 368	AW291698, AW449355, AA077663, and AI624833.
HELAR07	8792	953626	1 - 396	15-410	AA160979.
HEIAJ39	8793	705643	1 - 388	15 - 402	AA262593, AW055245, AA765419, Al268356, and Al919028.

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HEIA186	HEIAD76												٠					HEIAC62	HD/FMJ27				HDTL105	HDTLE28							

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HDTH102	8812	918728	1 - 468	15 - 482	AA441986, AA441987, AI051488, AL119444, AL119363, and AL134538.
HDTHG69	8813	754464	1 - 347	15-361	R05736, and AC001226.
HDTHE32	8814	851964	1 - 686	15 - 700	AW295293, AA005236, AW293791, AA564263, AA005235, and AL024494.
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HDTGT162	8816	742828	1 - 479	15 - 493	R10737, R10755, and AW206939.
HD/TGQ31	8817	697539	1 - 368	15 - 382	W57998, AA082443, A1979163, N50386, AA730309, AA282104, AL046056, and AF191338.
HDTGH10	8818	996896	1 - 399	15-413	AA968621, and AA911679.

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	1 - 219	1 - 2399	1 - 338
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	HDTAU14	HDTAT80	HDTAP53

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HDQPR06	8873	936658	1-815	15 - 829	AW276983, AW206541, AA418882, AI298636, AA290805, AA127225, AW368344, AI762472,

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HDQPP57	8874	926953	1-711	15 - 725	H80171, AA971126, and T80926.
HDQII08	8875	958252	1 - 456	15 - 470	AA010427.
HDQHU84	8876	782202	1 - 431	15 - 445	AW135842, Al185102, Al521545, N71336, Al374790, Al093509, H47630, AA580199, AI870424, AI935765, AA973252, AI538850, AI249946, AI909661, AI679550, AA514684, AW167422, AI687482, AW026905, AI933992, AI432570, AI684164, AI370322, AI440444, AI591407, AI687482, AW026905, AI933992, AI432570, AI684164, AI370322, AI440444, AI591407, AI570627, AW075921, AI817543, AI610318, AA243996, AW366372, AI963191, AI954721, AI242931, AI356929, AI270039, AI890391, AI570264, AW152415, AI089970, AI97746, AI679214, AI570857, AW089932, AI491904, AI624693, AI803740, AI682121, AI539800, AW166612, AW090102, AI446785, AI811912, AI830187, AW105665, AW167215, AW192461, AI439903, AI918370, AL046942, AI365256, AI687568, AW236186, AI922996, AI689614, AW170773, AW168503, AW163464, AI758988, AW0080085, AI68396, AI633967, AI689614, AW170773, AW168503, AV1639616, AW169234, AI582900, AW104117, AI539687, AL137298, U89906, AB007812, AB025103, AF114784, AL137471, AJ003118, AF182215, AL117648, X89102, AF194030, X59414, X84990, AL133559, AF109683, AL133113, E08517, E15582, A21103, AF114170, AL33062, AF061981, AF207750, A08915, AI8777, AR029580, U70981, I22020, AL137294, E01963, AF114170, AL133084, U75378, AL137268, AB0228451, AF1066945, AF000145, AF008145, AF008148, AP000083, S78214, AL137539, and N71386, AL117443, X62773,
НДОНО94	8877	793387	1 - 114	15 - 128	AA211827, and AA256750.
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IDQGW10	8879	963512	1 - 464	15 - 478	AW207166, R62452, AA570462, AW131259, U95742, and AC007216.
HDQGF78	8880	772952	1 - 546	15 - 560	R83407, and AC003086.
HDQFT10	8881	852587	1 - 685	15 - 699	AA864468, AI733017, and AI791415.
HDQFJ39	8882	852593	1 - 805	15 - 819	AA765774.
HDQFE93	8883	791724	1-1181	15 - 1195	T75255, H82395, R21558, H15880, F12818, Z42224, R13241, and F12817.
НДОЕХ85	8884	875473	1 - 495	15 - 509	AA002220, H63472, AA307783, AA478844, AI676034, AW364878, AI174745, AI682476, C04105, AA481986, U47346, R10969, T49897, and AL096740.
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HDQET29	8887	852618	1 - 459	15 - 473	T70748, AA584765, H86388, AI801563, AW167773, AA662629, AA572718, F27846, AL031427, Z97630, AL050318, AC007308, AL035423, AL109628, AC006121, D79206, AF053356, AC005255, AC006064, AC005592, AC005102, AJ239325, AC005971, AL035443, AL132777,

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15 - 445		15 - 441	15-361	15 - 395	15 - 388	15 - 593	15-919		15 - 593	15 - 632	15 - 607				15 - 569											15 - 895		15 - 400	15 - 466	15 - 776	15 - 360		
1-431		1 - 427	1 - 347	1 - 381	1 - 374	1-579	1 - 905		1 - 579	1 - 618	1 - 593				1 - 555											1 - 881		1 - 386	1 - 452	1 - 762	1 - 346		
934616 923120		744461	872237	690497	662917	923856	773487		923843	919597	764407				607575											674539		706647	493958	725029	934144		
8940	9,00	+		8944	8945	8946	8947		8948	8949	0568				8951										1	8952		8953	8954	8955	9568		
Н <u>р</u> РQH06 Н <u>р</u> РQF03	22000011	HDPQC63	HDPQA64	HDPPP29	HDPPB17	HDPOX03	HDPOT78		HDPOP03	HDPOC02	HDPN173				HDPND09											HDPNC22		HDPMV35	HDPMR81	HDPMH51	HDPMG46		

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	15 - 581	110-61	15 - 810				15-418	15 - 487	15 - 576	15-512		15 - 356	15 - 293	15 - 255	15 - 828	15 - 491	15 - 496	15 - 584	15 - 342
	1 - 567	1-59/	1 - 796				1 - 404	1 - 473	1 - 562	1 - 498		1 - 342	1 - 279	1 - 241	1 - 814	1 - 477	1 - 482	1 - 570	1 - 328
	849526	182001	973945				773469	923841	952966	716466		488326	887886	953001	933240	703959	771939	935198	764344
	8957	8938	8959				0968	8961	8962	8963		8964	8965	9968	2968	8968	6968	8970	1268
	HDPLV29	HUFLU84	HDPLT62				HDPKI78	HDPKA03	HDPJX07	НDРЉ44		HDPJK23	HDPJB19	HDPJA07	HDPIW95	HDPIV34	HDPIQ24	HDPIP06	HDPIL73

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1 - 330		
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8972		
HDPIL01		

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	15 - 361	15 - 445	15 - 570	15 - 423	15 - 430														15 - 473	15 - 327	15 - 509		15 - 390	15 - 508	15 - 514	15 - 462	15 - 728	15-410	15 - 467	15 - 468	15 - 369	15 - 467
	1 - 347	1 - 431	1 - 556	1 - 409	1 - 416														1 - 459	1-313	1 - 495		1 - 376	1 - 494	1 - 500	1 - 448	1 - 714	1 - 396	1 - 453	1 - 454	1 - 355	1 - 453
	881960	750575	729441	415993	728058		_												827350	706622	786078		711126	812336	704036	680099	964519	924010	973363	792256	921381	715232
	8973	8974	5268	9/68	2268														8678	6268	0868		8981	8982	8983	8984	8985	9868	8987	8868	6868	8990
	HDPIF78	HDPIF66	HDPIF54	HDPID88	HDPID53														HDPLA21	HDPHF37	HDPHA89		HDPHA40	HDPHA37	HDPHA34	HDPGX15	HDPGX10	HDPGX03	HDPGT96	HDPGS93	HDPGS45	HDPGS43

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15 - 461	15 - 468	15 - 227	15 - 466	15 - 456
1 - 447	1 - 454	1 - 213	1 - 452	1 - 442
909833	916352	468389	394693	416046
	8992	8993	8994	8995
HDPGS16	HDPGS01	HDPGR44	HDPGN15	HDPGN13

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	15 - 402	15-310	15 - 478	15 - 467
	1 - 388	1 - 296	1 - 464	1 - 453
	966923	722903	766618	765524
	9668	2668	8668	6668
	HDPGN11	HDPGJ49	HDPGI75	HDPGI74

	AA111917.	A1032747, A1769719, T79745, and AC004751.	AA225394, AA484596, AA484172, AL042753, AL042377, AI475660, AA464969, AW024544, AL045360, AA603359, AA808153. AL138455, AL134524, AA974503, AL043095, AL045360, AA603359, AA808153. AL138455, AL134524, AA974503, AL043095, AU17882, AL0402919, AA177033, W95924, AI440426, AL046110, AA947997, AI696846, AL042891, AA760555, AI860304, AC000581, AC000491, AF196779, AC005015, AC006571, AC006491, AL109627, AC005015, AC006571, AC006480, AL109671, AC005686, AL008719, AC005015, AC006571, AC005015, AC005210, AC005210, AC005210, AC005210, AC005210, AC005210, AC005210, AC005210, AC005210, AC005201, AC00501, AC005011,				T39977, and AL050337.	W72416, AA955016, and W76266.			
15-118	15-318	15 - 447	15 - 775	15 - 572	15-481	15 - 476	15 - 465	15-811	15 - 492	15 - 456	15 - 483
1 - 104	1 - 304	1 - 433	1 - 761	1 - 558	1 - 467	1 - 462	1 - 451	1 - 797	1 - 478	1 - 442	1 - 469
764397	916389	879489	709684	919306	486744	486737	796360	760327	486765	486738	486766
0006	9001	9002	9003	9004	9005	9006	2004	8006	6006	9010	9011
HDPGH73	HDPGG01	HDPGD28	HDPF270	HDPFW02	HDPFS71	HDPFS59	HDPFP96	HDPFP71	HDPFP14	HDPFN23	HDPFM28

HDPFL52 HDPFJ58	9012	486739 783957	1 - 469	15 - 483	AC007281. W05340.
HDPFD10	9014	964521	1 - 705	15-719	AA424163, and AI350948.
HDPFB03	\$106	936331	1 - 678	15 - 692	AI694127, AI299320, H70360, T98966, AA356015, AL045292, AA236204, AA131671, AW118104, W20295, H75919, N31893, R98291, AL117517, and AB023221.
Н DРDH08	9016	959303	1 - 344	15 - 358	AA082104, and AA134717.
HDPDD12	9017	970592	1 - 964	15 - 978	AI939431, AW020534, AI347310, AI371784, AI423466, AW150254, AW015338, AI361447,
					AI419452, AI088725, AI371086, AI418456, AI361506, AI419176, AI363726, AW176344,
					AL120300, AI424173, AW016674, AI288889, AF097450, AF146793, AF011568, AB002332, and
					AF000998.
НЪРЪС90	9018		1 - 641	15 - 655	T52186, and Z97630.
HDPCY47	6106	775412	1 - 402	15-416	W38488, N27984, AA649285, AA661773, AA298770, AA306867, H81350, AA100476,
					AA310276, AA417302, AA314322, AI815642, AA045597, D82429, R89903, AA130829, R83064,
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					N51146, AL122042, and AC007842.
HDPCP89	9020	786075	1 - 384	15 - 398	AA494163, T52475, and AC004148.
HDPCP64	9021	746527	1 - 320	15 - 334	AL022329, and AL022332.
HDPCP48	9022	721620	1 - 406	15 - 420	
HDPCP46	9023	719014	1 - 447	15 - 461	
HDPCP43	9024	715227	1 - 432	15 - 446	AI751151, AA418897, AA195596, AA491307, T54037, AA339713, AI751325, W79862, N41703,
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				_	AA216580, R72732, AA330733, AA227045, N36603, AA219518, W52126, AI336290, AA213719,
					H30288, K94642, AF112482, J04/94, AF036680, and AF112484.
HDPCP16	9025		1 - 408	15 - 422	
HDPC012	9056	852847	1 - 446	15 - 460	N23913, AI674840, AL079734, AI206841, AA502991, AA452887, AL119691, AA581247,
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	15-453	15 - 449	15 - 353	15 - 473	15 - 465
	1 - 439	1 - 435	1 - 339	1 - 459	1 - 451
	416035	764400	416028	852849	454740
	9027	9028	9029	9030	9031
	HDPCN88	HDPCN73	HDPCN33	HDPCM66	HDPCM62

APATOSTO, AAA13905, AAA13906, AAA43906, AAA4
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			15 - 406	15 - 675	15 - 501	15 - 417	15 - 1568	
		-	1 202	1 - 661	1 - 487	1 - 403 1 - 384	1 - 1554	
				783544	670893	719027		
			0030	9040	9041	9042	9044	
			HNDBEIK	IIDPBE85	HDPAX21	HDPAV46 HDPAP04	HDPAO68	

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HDPA064	9045	746550	1 - 445	15 - 459	T65324, AA575982, F09955, H88491, and AA947153.
HDPAN94	9046	901123	1 - 450	15 - 464	AA917335, Z20320, AL080176, AF144235, and A74743.
HDPAK09	9047	4941.70	1 - 437	15 - 451	AW168420, AI254913, F12561, AI382825, AA846952, F08248, AA572999, AI751216,
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					ALUSSAUS, UU/302, ACUU048U, ACUU489U, ALUZSS/S, ACUU40UU, ALUSI0S9, ACUUSSZ4, ACO06538, ACO07308, ACO05046, AI 117258, 191318, ACO06538, ACO07308, ACO05046, AI 117258, 191318, ACO06538, AI 078477, ACO16025
•					AC005512, U89337, AC007666, AC004797, Z97056, AC006966, AL022722, AC006039,
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	3				Z82198.
HDPA104	3048	881034	1 - 428	15 - 442	1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
HDPAE40	9050	856301	1 - 555	15 - 569	TOTODOR '1'UTODA A STAARS TOTODOK "A TOTODOR 1'UTODA A STAARS TOTODOK 1'UTODA A STAARS TOTODOK "A TOT
HDPAC90	9051	787796	1 - 806	15 - 820	4 A D 20808 HAR 260 A A 600610 A 1800670 W72575 A 1602860 A 1202015 A W220772 TS1767
	; }		}		T51608, and AA284050.
HDPAC78	9052	773570	1 - 226	15 - 240.	H82420, H82680, and AL078581.
HDPAB34	9053	281900	1 - 442	15-456	T94880.

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T84574, and AI192443.	AL037050, AW081941, AW406447, AA342901, AW088058, AA916039, AW074398, T71998,	AI299050, AI281903, AI689097, AA610491, AI054246, AI053816, N71724, F19258, AA095511,	AF177861, AA336362, AA377385, AW272758, AA501722, AA340003, AA30806, AA318652,	AI357551, AI565245, AI886176, AW236342, N22059, AI364809, AA491814, AI859251,	AA515435, AW301818, AJ919265, AL138265, AI287451, AA177061, F29989, AW440976,	A1053429, A1053672, A1274692, A1696962, AL040547, A1307426, A1284640, AA932099,	AI334443, T06556, AW340844, AW361583, AI358229, F36273, AA743956, AA553465,	A1963095, AW406755, AI200051, AI291124, AI365988, AI654588, AI291268, AW071063,	AI537030, AW339687, AI635272, AI133164, AA321311, AI216799, AI016000, AL041368,	AL138455, AA252763, AA652057, AA649642, AI264743, AI610159, AI469172, F24284,	AI690686, AI623132, AA745410, AL048616, AA654627, AA601892, AI537955, AA454925,	R13151, A1446601, A1801591, AA501784, F32705, AP000356, AC006004, AL121603, AC006064,	AC005215, AC004814, AC005940, AL034420, AC004491, Z73359, AP000245, AL049636,	AC006480, AF053356, AP000556, AC004894, AC002350, AP000552, AC005324, AL020993,	AL022324, AC002072, AC004895, AF045555, AF111168, AL121653, AC005619, AC002316,	AC003043, AC006285, AC004477, AC005800, U07562, AC005412, AC005081, AL121655,	AL121658, AC006001, AC005046, AC005099, AC005722, AL096701, Z84484, AF111169,	Z97054, AC007225, AC005920, AC007216, AC007899, AC005015, AF196779, AC007878,	AL035683, AC004966, AL033521, AL035460, U95742, AC004962, AC007242, AC006388,	AC000025, AC000083, AP000210, AP000132, AC004534, AC006111, AC004156, Z82214,	AL096791, AC007276, AC007371, AC007686, AC016025, AC005529, AC006312, AC008041,	AC004812, AL021453, AL033392, Z86061, AL132718, AF129756, AL031584, AC006153,	AL049779, AL021918, AC007707, AC005157, AC005037, AP000030, AP000558, AL023284,	AC005790, AL109967, AC007666, Z98742, U91323, AC007011, AP000248, AF001549,	AC004815, AC008134, AC005284, AC004890, AC006251, AC005066, AP000251, AP000557,	AC004964, AC005280, AC005696, AL024507, AC003982, AC006071, AC004841, AP000474,	AL049569, AC006211, AC008101, AC009330, AC007277, AC000078, AF196969, AL079340,	AC002070, AC000052, AL049830, AC004745, AC007450, AL024498, AC005180, AL133243,	AL031228, AC005666, AC003119, AC004150, AL035587, AL035090, AL035249, AC003104,	AC006195, U91326, AC005250, AL031005, Z98743, AC007537, Z98048, AJ251973, AC002432,	AC004776, AC007262, AC005694, AC004148, AC009516, AC004983, AC000159, AC005562,	Z84466, AC006057, Z82901, AL031848, AC005288, AC005625, AC005510, AC002369,	AL109802, U89337, X64467, AC007387, AL034451, AJ010770, AC007688, AB026898,	AL021391, AP000311, AC002312, AC004472, AC005755, AC004750, AL031777, U47924,	AL022323, Z93017, AC002525, Z82182, Z85987, AC004770, AP000692, AL035450, AF042090, AC004813, AL031230, AL132777, AC002115, AC004650, Z83844, AL008582, AC005911,
15-515	15 - 704																_							_											
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775417	546575																																		
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HDPAA80	HDPAA38																																		

HDLBEII	9026	966230	1-365	15 - 379	AC005324, AJ003147, AF017104, Z98946, AC007919, AC005821, AJ246003, AL117351, AC005041, AC0033041, AC007358, AL133355, AL035411, AC005919, AC008115, AL031255, AC005913, AF134726, AC006012, AC008079, AC005358, AC004694, AC009044, AC003102, AL109628, AL031053, U85195, AF109907, AL049766, AC006077, AL050341, AC005399, AC005331, AC005997, AL022165, AL078463, AC006323, AP000045, AC005242, Z99716, AC005295, AP000555, and AC004996. AA419383, AA411025, T03583, AA233509, AA969363, AA186898, AA485941, F12561, T65562, T59612, AW020599, AA989107, AA831152, F13808, AA745507, AW020088, AW022655,
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HDLAX52	9058	726241	1 - 819	15 - 833	AA033545, AW265437, and AA332853.
HDLAW58	9059	735559	1 - 869	15 - 883	AA461055, and Al371831.
HDLAG57	9061	565105	1 - 429	15 - 443	A1354333, AA715876, AI538236, AA828153, AW080910, AL049712, AC008079, AP000501,
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A A A	15 - 437	15 - 490	15 - 281	15 - 493 AZ	15 - 460 AI	15-430 AI	15 - 346 AZ	15 - 509 AZ	15 - 719 NS	15 - 461 AZ	15-510 AI) A(- AC	<u> </u>	<u>A</u>	- AC			15 - 579 N8	Z	Ž		15 - 582 NE	Ž	Ž, '	1		15 - 485 Al	15-611 H	15 - 648 N	
	1 - 423	1 - 476	1 - 267	1 - 479	1 - 446	1-416	1 - 332	1 - 495	1 - 705	1 - 447	1 - 496	•	•					1 - 457	1 - 565			İ	1 - 568				1 - 505	1 - 471	1 - 597	1 - 634	
	751699	799888	168661	736081	969401	951948	933818	852968	625387	966723	915420	•	_	·				974488	974462		_		974467				853002	964080	743064	974466	
	9062	9063	9064		9906	_	8906	6906	9070	9071	9072							9073	9074				9075				9076	9077	9078	9079	
	HDLAC67	HDLAC51	HDLAC30	HDLAA58	HDCBF12	HDCBD07	HDCAN06	HDCAG61	HDBAK09	HDBAA11	HCWWH01							HCWVA80	HCWUS57				HCWUS50				HCWUQ81	HCWUQ10	HCWUP62	HCWUM01	

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	15 - 693	15-410	15 - 444	15 - 622	15 - 547	15 - 441	15 - 347	15 - 354														_												
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	706530	722674	662486	919252	706523	927571	291888	757172								•								_										
	0806	9081	9082	9083	9084	9085	9806	2806																										
	HCWUL35	HCWUI49	HCWTP17	HCW/LL02	HCWSC38	HCWLA43	HCWKT71	HCWKS70																										

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	15 - 333	15 - 383	15 - 285																											15 - 328		
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	8806	6806	0606																											9091		
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HCWKR51	9092	725027	1 - 329	15 - 343	TOTAL CALEBOAR STORMS OF ONE STORMS STORMS STORMS STORMS
HCWKR14	9093	658421	1 - 413	15 - 427	AI3/6239, AI88/314, AI22438/, AI652810, CI5060, AI85//56, AI45/313, AI63418/,
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					AA811741, AA743968, AA218720, AA084609, AW082470, AA368616, AA604865, AA548886,
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	9106	6107	8016	9109							3	9110	9111		•			9112					9113										
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					Z86061, AC006480, AC004878, AL035460, AC006241, AC006039, AC002389, AC005071,
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HCWAB25	5 9143	575270	1-151	15 - 165	AA630672, AW148507, H18914, AL047429, AA640410, AA640430, AA127426, AI292257,
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					AA018207, AA714011, H25921, AW410354, AA101689, AI341571, AA629872, AA507612,
					AI583466, AW303096, AW268291, AW274191, AI802801, AA504049, AA507187, AI783549,
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-					ACCOVIZZO, ACCOCASOS, DOISOS, ALOSOVOI, ACCIOUSZS, ASOCOSIAI, ALDOVISOZ, ACCOSIZS, APPONOSIA APPONINGO AL 1212245 ACCOCASOR ACCOCASOR ACCOCASOR ACCOCASOR ACCOCASOR
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IICVAC70	9149	507284	1 - 801	15 - 815	AA196390, AA196468, AA507837, AI537538, AA845293, AI051037, AI500454, AL038936, F27015, AI446601, F23315, AA772016, AI982634, AI954525, AI744949, AI708005, AA064938, AA229444, AW157005, AC001228, AC005996, AC005210, AC007436, AJ003147, AC005399,
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					A1921619, 298474, AA708651, R17935, AA076130, AA743299, NS5076, H65911, AA467774,
					T59438, AA250776, AA856817, R48980, AI348467, AA302763, AA115865, AA488687.
· ve					AA363957, N30016, AI569700, AA507993, AA524821, AI924954, AI908424, T86603, R53995,
					AI251024, AA065155, AA247477, AI453160, AI564294, AC007671, D83737, AC006962,
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					AC009294, D84394, Z69366, AP000228, AP000140, AP000088, AC006479, AF147446,
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	15 - 340	15 - 289	15 - 321	15 - 332	15-418	15 - 223	15 - 333	15-655	15 - 431	15 - 737	15 - 388	15 - 424	15 - 395	15 - 369	15 - 418	15 - 327							
	1 - 326	1 - 275	1 - 307	1 - 318	1 - 404	1 - 209	1 - 319	1-641	1-417	1 - 723	1 - 374	1 - 410	1 - 381	1 - 355	1 - 404	1 - 313							
	861969	529691	529692	529694	869625	529695	879694	934889	846648	765645	657297	964727	820008	730739	792756	765641							
	9151	9152	9153	9154	9155	9126	9157	8516	9159	9160	9161	9162	9163	9164	9165	9166							
	HCVAC47	HCVAC45	HCVAC38	HCVAC33	HCVAC26	HCVAC21	HCVAC19	HCUIO06	HCUIM28	HCUIK74	HCUHY13	HCUHY10	HCUHU61	HCUHU54	HCUHT92	HCUHT74							

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	15-250	15-119	15 - 244					_																							
	1 - 236	1 - 105	1 - 230									_																			
	575794	681874	657295																												
	1916	9168	6916				_																								
	HCUHT37	HCUHT26	HCUHT13																												

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	15 - 554	15 - 499	15 - 1071	15-437
	1 - 540	1 - 485	1 - 1057	1 - 423
	924494	690027	879921	967142
	9170	9172	9173	9174
·	HCUHT03	нсино27	нсино13	HCUHK11

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959567	551796	796434								
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HCUDN82 HCUDN71	9254 9255 9256	779978 862116 959679	1 - 429 1 - 381 1 - 395	15 - 443 15 - 395 15 - 409	AI823714, AI807753, AA084901, AA284234, AI312027, AI436666, AW001163, AA372073, AF076931, AF099739, AB016869, and AB019245. T63603. AA278703, and AW292815. AI629017.
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HCUDD07	9259	953588 953588	1 - 346	15 - 433	AA086020. AL117375, AL133163, AC006512, AC006241, AC005225, AL050348, AC007240, AC006450, AL117375, AL1375, AL133163, AC006512, AC006491, AC004973, AC005513, AC004912, AC005519, AC005003, U07563, Z85987, AP001054, AP000343, AC008134, L78810, AC005006, AL109984, AC005697, AC005300, AC004882, AC005399, AL133371, U91321, Z97054, AC006946, AB001523, AC005081, AC006465, and Z84466.
HCUCK58	9260	850117	1 - 253	15 - 267	N78946.
HCUCI23	9262	676276	1 - 196	15-204	AA021595.
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HCUCG40	9264	800153	1 - 141	15 - 155	
HCUCG07	9265	953823	1-287	15 - 301	AA807580.
HCUCD83	9266	781346	1-411	15 - 425	N29857.
HCUBX45	9268	850122	1 - 497	15-511	

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1-431	1 - 312	1 - 386	1 - 137	1 - 56	1 - 397	1 - 370
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6976	9270	9271	9272	9273	9274	9275
HCUBV91	HCUBT35	HCUBS07	HCUBP94	HCUBP82	HCUBP48	HCUBP34

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HCUBO67	9277	526489	1 - 261	15 - 275	AA533534, AA569471, AI434695, AW407578, T60681, AI801482, AA760839, AW408047, AA630672, AA502991, AI932599, F11718, AA224889, AW328202, AL048925, AI143051, F00564, AW270258, AW002350, AA829166, AC008115, AC005666, AC005800, AP000212, AP000134, AP000251, L44140, AF001548, AP00030, U95742, AL049872, AC007216, AC002312, Z68870, AC006042, AC005839, AC005231, AC002430, AC0064517, AC007421, AC002306, AC007384, AP000047, AC005482, AC005520, AC005632, AC016025, AL096767, AF053356, AC005808, AC003665, U63721, AL03216, Z94056, AC005221, AL035685, AC004967, AP000115, AC004878, AP000553, AC007934, AC004216, Z54334, AL049539, AC004967, AL09798, AL021155, AL035659, AL137100, AC004587, AP000152, AC004885, AC005667, AL035400, AC004813, AC005886, AP000252, AC005005, AC004586, AC006885, AF196779, AC004458, AC002300, U91323, AC004707, AL022302, AC006254, AC006285.

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	1 - 442 1 - 638 1 - 455
	577294 722190 968223
	9278 9279 9280
	HCUBM73 HCUBL10 HCUBL10

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_	9285 723175	1-	196	15 - 210	T58148, and AW059603.
HCUBE22	86 409828	1 - 714	14	15 - 728	R10475, AA682937, AA601674, R31425, AA689285, AI783581, AW190781, R06346, AA922351, B10346, AA607745, A1567833, AW008080, A134536, A1310873, A1180689, B21380
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	1 - 596	1-514	1 - 325	1-414	1 - 204	1 - 224	1 - 129	1-151	1 - 509	1 - 345	1 - 154
	752247	792515	917266	924897	706426	796010	574275	728364	887470	702677	968229
	9287	9288	+	9290	\vdash	┢	9293		9295	9296	9297
·	нсивв68	HCUBA93	HCUAZ01	HCUAW03	HCUAP34	HCUAN95	HCUAN89	HCUAN53	HCUAM65	HCUAL33	HCUAG10

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15 - 419	15 - 599	15 - 639
1 - 405	1 - 585	1 - 625
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9299	9300	9301
HCUAF01	HCUAA92	HCFOP90

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HCFMH46	9322	719113	1 - 342	15 - 356	AW239548, AW373232, and AA325059.
HCFLY85	9323	909182	1 - 508	15 - 522	AI340241, AI740877, AA262686, AI949440, AA011068, AA011069, AA281890, AA283910, AA527101, AA250824, AI340240, and AA255646.
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HCFLY08	9325	959620	1 - 926	15 - 940	AA485570, AA485409, AI458623, R22065, AA639708, AL134139, AL031678, AC004491, AC004985, and Z99716.
HCFLV30	9326	692781	1 - 448	15 - 462	T89518, N91297, R05742, N78236, AA281577, AA505098, and AA282794.

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	1 - 323 1 - 993		1 - 683	1 - 806	1 - 393	1 - 341		
	614915 847853		772000	952843	771975	746398		
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	HBJLE70		HBJK177	HBJKG07 HRJKF02	+			

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HBJKD40	9449	710931	1 - 351	15 - 365	
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HBJKA01	9451	916077	1 - 528	15 - 542	A1445151.
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НВ ЈНU20	9468	907575	1 - 544	15 - 558	AA179630, and AA176615.
HBJHO44	9469	882040	1 - 409	15 - 423	R96772.
HBJHI96	9470	828996	1 - 424	15 - 438	AA461038, AW151827, AI753072, AI042290, AI377120, AI364168, W68727, N30339, AI754813,
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					AW338200, AW291880, AW33/350, AI913188, AI243881, AA291380, H/3392, AA/02481,

			W85789, AA693974, AI032087, and W85723.			AI018729, AI017735, T94096, AA788760, AC004878, AC006014, AC005488, AC005071, AC006480, AF030453, and AC005088.	R96956,	N92225.	H02200.	R92654, R92648, and N59587.	D58283, D80195, D80227, D80391, D80166, D59859, D80043, D59502, D51423, D59619,	D80210, D51799, D80240, D80253, D80196, D80269, D80038, D80188, D80212, D59275,	D80193, D57483, D59927, D80219, D81030, D80022, D50979, D59889, D59610, D80366,	D80378, D80045, D59787, C14429, D50995, D80164, D81026, D80241, D80024, T03269,	C15076, C14389, C75259, C14014, C14331, D59467, D51060, AW178893, AA305409, D80134,	F13647, D51250, D80268, D58253, D80949, AA305578, D80168, C14227, D51079, D81111,	D51022, AW177440, AW179328, D80522, D52291, AW178775, AW378532, D59695,	AW352158, AW377671, Z21582, AI910186, AW369651, AI905856, D80251, AA514188, D80248,	AW178762, AA514186, D51097, AA285331, C05695, C14298, AW177501, AW177511, D80064,	D80133, AW360811, C14407, AW352117, AW176467, AW378540, AW375405, AW360834,	D80132, AW366296, AW360844, AW360817, AW375406, AW378534, AW179332, AW377672,	AW179023, AW178905, AW179220, D80302, AI557751, AW352171, D80439, AW377676,	AW178906, AW352170, AW17731, AW178907, AW179019, AW179024, D59373, D80247,	D80014, AW177505, AW179020, AW360841, AW178909, AW177456, AW179329, AW178980,	AW177733, AW378528, AW178908, AW178754, AW179018, AW352174, T11417, D51103,	T03116, AW179004, AW179012, AW178914, AW378525, D80157, AW367967, AW17722,	AW177728, T02974, D51759, AW179009, AW178774, AW178911, AW378543, AW352163,	D58246, C06015, D59503, D80258, AW178983, AW352120, AW178781, T48593, D51213,	AW378539, D59627, AI557774, D58101, AI535850, AA809122, AW177723, D59653,	AW177508, D45260, C14975, AI525923, AW378533, AW367950, H67866, D59317, H67854,	C03092, D45273, AW177497, AJ535686, AW178986, N66429, AW177734, AJ525917, C14973,	C14344, D59474, AI525227, D51221, D59551, D60010, AA033512, AI525920, AA514184,	D60214, C14957, C14046, C16955, T03048, AI525242, AI525235, AI525912, AI525925,	AI535961, AW378542, AI525215, C05763, Z33452, AI525237, AI525222, D31458, AJ132110,	A62300, A84916, A62298, X67155, A67220, D89785, A78862, D26022, A25909, Y1/188,
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1 - 424	1 - 105	1 - 507	1 - 598	1 - 96	1 - 255	1 - 383	1-415	1 - 422	1 - 386	1 - 624	1 - 807																								
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HBJCJ82	9509	779645	1 - 467	15 - 481	
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HBJA055	9514	732436	1 - 414	15 - 428	T81741, AA778192, W85729, and AL132777.
HBJAJ49	9515	723164	1 - 427	15 - 441	N71474,
HBJAI50	9196	724619	1 - 490	15 - 504	A1089358, AA056686, A1089564, A1086521, A1669927, A1817821, AA056582, AA847148,
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HBJAH27	9517	682941	1 - 239	15 - 253	
HBJAE32	9518	699695	1 - 395	15 - 409	W88646, AI368895, and Z39215.
HBJAB77	6156	772422	1 - 491	15 - 505	H53026.
HBJAB49	9520	722195	1 - 874	15 - 888	AI479236, AI935423, AA255972, and AA243356.
HBJAB28	9521	847966	1 - 486	15 - 500	Z28561.
IIBJAB15	9522	660552	1 - 395	15 - 409	AA429379.
HBDAE47	9523	720008	1 - 572	15 - 586	AA252703, AI760942, and AA565583.
HBDAD64	9524	864344	1 - 534	15 - 548	H04563, R80434, AA531428, Z82215, and Z95114.
HBDAD16	9525	\neg	1 - 375	15 - 389	AI810593, and T89570.
HBCC010	9526	963157	1 - 564	15 - 578	AA701314, and AA004793.

H59931, H67432, Z70280, AC003100, AC006998, AC003072, AC005154, AL022154, and AC003034.	A1672457, AA810483, AI269914, AI949892, AI198001, W86511, AW237699, AI863804, and AI525426.	AI078128, AI871101, AI274339, AA035467, AA419038, AI479404, AI677732, AW190726, AI198435, AA031617, AI283200, AI983969, AA411122, AI093316, AI361637, AI769275, AI560217, AA961077, AI291805, AI291474, AA915909, AA035466, AA046943, AI857306, AA661657, AA427407, AA725194, AA423792, AW070742, AW051192, AI275083, AW293787, AW452102, AA250784, AA031475, N92812, H91759, AW149476, H91665, and AA514348.		AA280776, AL137689, AF111173, and AF100172.	AA464030, AA434281, AA410674, D78694, A1860751, A1628190, AA568168, A1097605, A1199027, D78920, AA227411, A1200649, A1199599, A1935345, AA525327, A1475176, A1910441, A1923458, A1247190, A1687257, A1923451, A1541551, A1699991, A1653151, and AW438934.	H40594.	AA677024, and AI022041.	AI566483, AA417101, and AL021368.	AA486018, AA005417, H56995, AL031666, and AF144233.	AA419501, AA969163, AA157911, AA410996, AW452982, AW339854, and H14554		AA947361, Z97054, AL049694, AC005484, AC004253, AC005288, AC004019, AL049757,	AC004584, AC006441, AC009247, AC007395, AC004216, AC004895, AL022316, U91318,	U95742, AL049766, AL009181, AL023553, AC016830, AC002350, AC007216, AC000052,	Z85986, AC007676, AC005562, AC016027, AC005822, AL031848, AL050318, AL031662,	AC006088, AC003950, AC005086, AC004099, AB001523, AP000694, AC004841, AC010205,	AC004765, AC005280, AC004813, AL109984, AC005409, AC003101, AC004678, AC004812,	AP000514, AL034429, AC004874, AL031602, AC006487, AJ009610, AC004596, AL136295,	AC002314, AC007227, AL117694, AP001060, AC004125, AL031587, AL133246, AC005057,	AF001548, AP000512, AC005004, AC005088, Z83844, AL031311, U91323, AC007686,	AF001549, AL022165, AF030453, Z98941, Z84469, AC004383, AC005914, Z93016, AC005516,	AL049569, AC006059, Z93017, Z83847, AC005837, AC004913, AF217403, AC005015,	AC004662, AC005081, AC007193, AC008039, AC003982, AC005058, AC006130, M89651,	AC008372, AF043945, AC005696, AL096701, AL024474, AC002312, AL031276, AC0004531,	AL009183, AL049776, AL049780, AC005821, AC003046, AF134726, AL096712, AL031985,	AL049758, AL035249, AL023807, AP000501, Z93241, AL133382, AC005940, AC010077, AC008101 AL035086 AC005430 AC002425 Z86090 AC005082 AC007899, AC005049.
15 - 582	15 - 593	15 - 604	15 - 529	15 - 581	15 - 1009	15 - 539	15 - 602	15 - 531	15 - 772	15 - 909	15-363															
1 - 568	1 - 579	1 - 590	1 - 515	1 - 567	1 - 995	1 - 525	1 - 588	1 - 517	1 - 758	1 - 895	1 - 349		_								<u>.</u>					
930956	698596	938319	975256	958151	952057	934110	926761	969578	848322	848325	573919															
9527	9528	9529	9530	9531	9532	9533	9534	9535	9536	9537	9538															
HBCCJ05	нвссе11	НВССD06	HBCCB51	HBCBR08	HBCBN51	IIBCBN06	HBCBG04	HBCBF12	HBCBE57	HBCBB22	HBCAW52															

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	15 - 337	- 278	-376	15-432																						
	1 - 323	1 - 264	1 - 362	1-418																						
	968195	699489	783431	864356			_																			
	9539	9540	9541	9542																						
	HBCAT10	HBCAS32	HBCAQ85	HBCAQ42																			<u> </u>			

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	15 - 503	15 - 836
	1 - 489	1-1185
	738423	879377
	9543	9545
	HASCG58	HASAR74

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AF109901, AC00000 AF009810, AC00412 AC006333, AL10995 Z98200, AC006530, AC002416, AC00459 AI863136, N71484, A	ALUSUSSO, AF 15472 AF109907, AC00608	CENCIST OCCUSO IN	AL031431, AC00541	AC000385, AL13629 AL031431, AC00541	AL035420, Z85986, AC000385, AL13629 AL031431, AC00541	AC004386, AC00588 AL035420, Z85986, A AC000385, AL13629 AL031431, AC00541	AC004859, U78027, AC004386, AC00588 AL035420, Z85986, A AC000385, AL13629 AL031431, AC00541	AC004707, AC00235 AC004859, U78027, AC004386, AC00588 AL035420, Z85986, A AC000385, AL13629 AL031431, AC00541	AL135783, AC00449 AC004707, AC00235 AC004859, U78027, AC004386, AC00588 AL035420, Z85986, AC00531 AC000385, AL13629	AP000086, AL07863 AL135783, AC00449 AC004707, AC00235 AC004859, U78027, AC004386, AC00588 AL035420, Z85986, A AC000385, AL13629 AL031431, AC00541	AC003004, AL00871 AP000086, AL07863 AL135783, AC00449 AC004707, AC00235 AC004859, U78027, AC004386, AC00588 AL035420, Z85986, AC00531
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1-470											
HASAC10 9546 968746 1-470		_									
9546											
AC10							•			·	

TABLE 4

Code	Description	Tissue	Organ	Cell Line	Disease	Vector
AR022	a_Heart	a_Heart	Otgan	Cen Eme	Discase	Vector
AR023	a_Liver	a_Liver		-		 -
AR024	a_mammary gland	a_mammary gland				
AR025	a Prostate	a_Prostate		·		
AR026	a_small intestine	a_small intestine	 	 		
AR027	a_Stomach	a_Stomach				
AR028	Blood B cells	Blood B cells	· · · · · · · · · · · · · · · · · · ·	·		
AR029	Blood B cells activated	Blood B cells				
}	}	activated	})		}
AR030	Blood B cells resting	Blood B cells resting				
AR031	Blood T cells activated	Blood T cells activated				
AR032	Blood T cells resting	Blood T cells resting				
AR033	brain	brain		·		
AR034	breast	breast	·	1		
AR035	breast cancer	breast cancer				
AR036	Cell Line CAOV3	Cell Line CAOV3				
AR037	cell line PA-1	cell line PA-1				
AR038	cell line transformed	cell line transformed				
AR039	colon	colon				
AR040	colon (9808co65R)	colon (9808co65R)				
AR041	colon (9809co15)	colon (9809co15)				
AR042	colon cancer	colon cancer				
AR043	colon cancer (9808co64R)	colon cancer (9808co64R)				
AR044	colon cancer 9809co14	colon cancer 9809co14				
AR045	com clone 5	com clone 5				
AR046	corn clone 6	com clone 6				
AR047	corn clone2	corn clone2				
AR048	corn clone3	corn clone3		 		
AR049	Corn Clone4	Corn Clone4		<u> </u>		
AR050	Donor II B Cells 24hrs	Donor II B Cells 24hrs				
AR051	Donor II B Cells 72hrs	Donor II B Cells 72hrs				
AR052	Donor II B-Cells 24 hrs.	Donor II B-Cells 24 hrs.				
AR053	Donor II B-Cells 72hrs	Donor II B-Cells 72hrs				
AR054	Donor II Resting B Cells	Donor II Resting B Cells				
AR055	Heart	Heart		1		
AR056	Human Lung (clonetech)	Human Lung (clonetech)				
AR057	Human Mammary (clontech)	Human Mammary (clontech)				
AR058	Human Thymus (clonetech)	Human Thymus (clonetech)				
AR059	Jurkat (unstimulated)	Jurkat				
ADOCO	Vidnou	(unstimulated) Kidney		 		
AR060	Kidney	Liver		1		
AR061	Liver	Liver (Clontech)		 		
AR062	Liver (Clontech)	Lymphocytes		 		
AR063	Lymphocytes chronic lymphocytic leukaemia	chronic lymphocytic leukaemia				

AR064	Lymphocytes diffuse large	Lymphocytes				
AK004	B cell lymphoma	diffuse large B cell				
	B cell tymphoma	lymphoma			İ	
10066	1	Lymphocytes			·	
AR065	Lymphocytes follicular	follicular lymphoma			1	
1777	lymphoma	normal breast			 	
AR066	normal breast				 	
AR067	Normal Ovarian	Normal Ovarian			\	
	(4004901)	(4004901)			 	
AR068	Normal Ovary 9508G045	Normal Ovary			1	
		9508G045			ļ	
AR069	Normal Ovary 9701G208	Normal Ovary			1	
		9701G208			ļ	
AR070	Normal Ovary 9806G005	Normal Ovary			Ì	
		9806G005				
AR071	Ovarian Cancer	Ovarian Cancer				
AR072	Ovarian Cancer	Ovarian Cancer	i	1	l	1
	(9702G001)	(9702G001)				<u> </u>
AR073	Ovarian Cancer	Ovarian Cancer			j	1
	(9707G029)	(9707G029)				ļ
AR074	Ovarian Cancer	Ovarian Cancer			l	Į.
	(9804G011)	(9804G011)			L	<u> </u>
AR075	Ovarian Cancer	Ovarian Cancer			1	
	(9806G019)	(9806G019)			<u> </u>	
AR076	Ovarian Cancer	Ovarian Cancer		1		
	(9807G017)	(9807G017)				
AR077	Ovarian Cancer	Ovarian Cancer				
	(9809G001)	(9809G001)			ļ	
AR078	ovarian cancer 15799	ovarian cancer			i	
'		15799				
AR079	Ovarian Cancer	Ovarian Cancer				
	17717AID	17717AID			ļ	ļ
AR080	Ovarian Cancer	Ovarian Cancer			1	
	4004664B1	4004664B1				
AR081	Ovarian Cancer	Ovarian Cancer				
	4005315A1	4005315A1				
AR082	ovarian cancer 94127303	ovarian cancer	_		ł]
		94127303				ļ
AR083	Ovarian Cancer 96069304	Ovarian Cancer			İ	ļ
		96069304				
AR084	Ovarian Cancer 9707G029	Ovarian Cancer	_		ļ	ł
		9707G029			<u> </u>	
AR085	Ovarian Cancer 9807G045	Ovarian Cancer			1	
		9807G045			ļ	<u> </u>
AR086	ovarian cancer 9809G001	ovarian cancer			1	
		9809G001			ļ	ļ
AR087	Ovarian Cancer	Ovarian Cancer				
	9905C032RC	9905C032RC			ļ	ļ
AR088	Ovarian cancer 9907 C00	Ovarian cancer 9907			1	}
	3rd	C00 3rd			ļ	<u> </u>
AR089	Prostate	Prostate			ļ	
AR090	Prostate (clonetech)	Prostate (clonetech)			<u> </u>	<u> </u>
AR091	prostate cancer	prostate cancer				
AR092	prostate cancer #15176	prostate cancer				1
		#15176	L			<u> </u>
AR093	prostate cancer #15509	prostate cancer				
		#15509		<u> </u>	<u> </u>	<u> </u>
AR094	prostate cancer #15673	prostate cancer			1	T
ふんりブサ	prostate cancer #15075	#15673	}	}	1	l
AR095	Small Intestine (Clontech)	Small Intestine	 			
びどのとい	Small fillestine (Ciontech)	(Clontech)			1	
AR096	Spleen	Spleen				
AR090 AR097	Thymus T cells activated	Thymus T cells			 	
み K U サ /	I THYTHUS I COHS ACHVAICH	L Injinus i cons	L		·	

		activated	l			T
AR098	Thymus T cells resting	Thymus T cells resting				
AR099	Tonsil	Tonsil				
AR100	Tonsil geminal center centroblast	Tonsil geminal center centroblast				
AR101	Tonsil germinal center B	Tonsil germinal center B cell				
AR102	Tonsil lymph node	Tonsil lymph node				
AR103	Tonsil memory B cell	Tonsil memory B				
AR104	Whole Brain	Whole Brain				
AR 105	Xenograft ES-2	Xenograft ES-2				
AR106	Xenograft SW626	Xenograft SW626				<u> </u>
H0004	Human Adult Spleen	Human Adult Spleen	Spleen			Uni-ZAP XR
H0013	Human 8 Week Whole Embryo	Human 8 Week Old Embryo	Embryo			Uni-ZAP XR
H0022	Jurkat Cells	Jurkat T-Cell Line				Lambda ZAP II
H0025	Human Adult Lymph Node	Human Adult Lymph Node	Lymph Node			Lambda ZAP II
H0031	Human Placenta	Human Placenta	Placenta		-	Uni-ZAP XR
H0050	Human Fetal Heart	Human Fetal Heart	Heart			Uni-ZAP XR
H0057	Human Fetal Spleen					Uni-ZAP XR
H0058	Human Thymus Tumor	Human Thymus Tumor	Thymus		disease	Lambda ZAP II
H0059	Human Uterine Cancer	Human Uterine Cancer	Uterus		disease	Lambda ZAP II
H0060	Human Macrophage	Human Macrophage	Blood	Cell Line		pBluescript
H0061	Human Macrophage	Human Macrophage	Blood	Cell Line		pBluescript
H0062	Human Thymus	Human Thymus	Thymus			Uni-ZAP XR
H0063	Human Thymus	Human Thymus	Thymus			Uni-ZAP XR
H0069	Human Activated T-Cells	Activated T-Cells	Blood	Cell Line		Uni-ZAP XR
H0073	Human Leiomyeloid Carcinoma	Human Leiomyeloid Carcinoma	Muscle		disease	Uni-ZAP XR
H0074	Human Platelets	Human Platelets	Blood	Cell Linc		Uni-ZAP XR
H0075	Human Activated T-Cells (II)	Activated T-Cells	Blood	Cell Line		Uni-ZAP XR
H0083	HUMAN JURKAT MEMBRANE BOUND POLYSOMES	Jurkat Cells				Uni-ZAP XR
H0087	Human Thymus	Human Thymus				pBluescript
H0090	Human T-Cell Lymphoma	T-Cell Lymphoma	T-Cell		disease	Uni-ZAP XR
H0108	Human Adult Lymph Node, subtracted	Human Adult Lymph Node	Lymph Node			Uni-ZAP XR
H0109	Human Macrophage, subtracted	Macrophage	Blood	Cell Line		pBluescript
H0116	Human Thymus Tumor, subtracted	Human Thymus Tumor	Thymus			pBluescript
H0128	Jurkat cells, thiouridine activated	Jurkat Cells				Uni-ZAP XR
H0129	Jurkat cells, thiouridine activated, fract II	Jurkat Cells				Uni-ZAP XR

H0134	Raji Cells, cyclohexamide treated	Cyclohexamide Treated Cem, Jurkat, Raji, and Supt	Blood	Cell Line	Uni-ZAP XR
H0139	Activated T-Cells, 4 hrs.	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0140	Activated T-Cells, 8 hrs.	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0141	Activated T-Cells, 12 hrs.	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0158	Activated T-Cells, 4 hrs., ligation 2	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0159	Activated T-Cells, 8 hrs., ligation 2	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0160	Activated T-Cells, 12 hrs., ligation 2	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0161	Activated T-Cells, 24 hrs., ligation 2	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0167	Activated T-Cells, 24 hrs.	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0179	Human Neutrophil	Human Neutrophil	Blood	Cell Line	Uni-ZAP XR
H0185	Activated T-Cell labeled with 4-thioluri	T-Cells	Blood	Cell Line	Lambda ZAP II
H0186	Activated T-Cell	T-Cells	Blood	Cell Line	Lambda ZAP II
H0187	Resting T-Cell	T-Cells	Blood	Cell Line	Lambda ZAP II
H0189	Human Resting Macrophage	Human Macrophage/Monoc ytes	Blood	Cell Line	Uni-ZAP XR
H0191	Human Activated Macrophage (LPS), thiour	Human Macrophage/Monoc ytes	Blood	Cell Line	Uni-ZAP XR
H0202	Jurkat Cells, cyclohexamide treated, subtraction	Cyclohexamide Treated Cem, Jurkat, Raji, and Supt	Blood	Cell Line	Uni-ZAP XR
H0203	Jurkat Cells, cyclohexamide treated, dif	Cyclohexamide Treated Cem, Jurkat, Raji, and Supt	Blood	Cell Line	Uni-ZAP XR
H0218	Activated T-Cells, Ohrs, subtracted	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0219	Activated T-Cells, Ohrs, differentially expressed	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0220	Activated T-Cells, 4 hrs, subtracted	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0221	Activated T-Cells, 4 hrs, differentially expressed	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0222	Activated T-Cells, 8 hrs, subtracted	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0223	Activated T-Cells, 8 hrs, differentially expressed	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0224	Activated T-Cells, 12 hrs, subtracted	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0225	Activated T-Cells, 12hrs, differentially expressed	Activated T-Cells	Blood	Cell Line	Uni-ZAP XR
H0250	Human Activated Monocytes	Human Monocytes			Uni-ZAP XR
H0253	Human adult testis, large inserts	Human Adult Testis	Testis		Uni-ZAP XR
H0254	Breast Lymph node cDNA library	Breast Lymph Node	Lymph Node		Uni-7.AP XR
H0255	breast lymph node CDNA	Breast Lymph Node	Lymph Node		Lambda

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	library					ZAPII
H0261	H. cerebellum, Enzyme subtracted	Human Cerebellum	Brain			Uni-ZAP XR
H0264	human tonsils	Human Tonsil	Tonsil			Uni-ZAP XR
H0265	Activated T-Cell (12hs)/Thiouridine labelledEco	T-Cells	Blood	Cell Line		Uni-ZAP XR
H0271	Human Neutrophil, Activated	Human Neutrophil - Activated	Blood	Cell Line		Uni-ZAP XR
H0272	HUMAN TONSILS, FRACTION 2	Human Tonsil	Tonsil			Uni-ZAP XR
H0274	Human Adult Spleen, fractionII	Human Adult Spleen	Spleen			Uni-ZAP XR
H0300	CD34 positive cells (Cord Blood)	CD34 Positive Cells	Cord Blood			ZAP Express
H0305	CD34 positive cells (Cord Blood)	CD34 Positive Cells	Cord Blood			ZAP Express
Н0306	CD34 depleted Buffy Coat (Cord Blood)	CD34 Depleted Buffy Coat (Cord Blood)	Cord Blood			ZAP Express
H0318	HUMAN B CELL LYMPHOMA	Human B Cell Lymphoma	Lymph Node		disease	Uni-ZAP XR
H0341	Bone Marrow Cell Line (RS4;11)	Bone Marrow Cell Line RS4:11	Bone Marrow	Cell Line		Uni-ZAP XR
H0354	Human Leukocytes	Human Leukocytes	Blood	Cell Line		pCMVSport
H0369	H. Atrophic Endometrium	Atrophic Endometrium and myometrium				Uni-ZAP XR
H0370	H. Lymph node breast Cancer	Lymph node with Met. Breast Cancer			disease	Uni-ZAP XR
H0376	Human Spleen	Human Adult Spleen	Spleen			pCMVSport
H0393	Fetal Liver, subtraction II	Human Fetal Liver	Liver			pBluescript
H0402	CD34 depleted Buffy Coat (Cord Blood), re-excision	CD34 Depleted Buffy Coat (Cord Blood)	Cord Blood			ZAP Express
H0416	Hurnan Neutrophils, Activated, re-excision	Human Neutrophil - Activated	Blood	Cell Line		pBluescript
H0421	Human Bone Marrow, re- excision	Bone Marrow				pBluescript
H0422	T-Cell PHA 16 hrs	T-Cells	Blood	Cell Line		pSport1
H0423	T-Cell PHA 24 hrs	T-Cells	Blood	Cell Line		pSport1
H0436	Resting T-Cell Library, Il	T-Cells	Blood	Cell Line		pSport1
H0439	Human Eosinophils	Eosinophils				pBluescript
H0444	Spleen metastic melanoma	Spleen, Metastic malignant melanoma	Spleen		disease	pSport1
H0445	Spleen, Chronic lymphocytic leukemia	Human Spleen, CLL	Spleen		disease	pSport1
H0457	Human Eosinophils	Human Eosinophils				pSport1
H0477	Human Tonsil, Lib 3	Human Tonsil	Tonsil			pSport1
H0478	Salivary Gland, Lib 2	Human Salivary Gland	Salivary gland			pSport1
H0485	Hodgkin"s Lymphoma I	Hodgkin"s Lymphoma I			disease	PCMVSport 2.0
H0486	Hodgkin"s Lymphoma ll	Hodgkin"s Lymphoma II			disease	pCMVSport 2.0
H0487	Human Tonsils, lib I	Human Tonsils				pCMVSport 2.0
H0488	Human Tonsils, Lib 2	Human Tonsils				pCMVSport

H0494	Keratinocyte	Keratinocyte				2.0 pCMVSport
TIOCOC			Colon	ļ		2.0 pSport1
H0506	Ulcerative Colitis	Colon Human Liver.	Liver	 		
H0510	Human Liver, normal	normal, Patient #8	Liver			pCMVSport 3.0
H0518	pBMC stimulated w/ poly	pBMC stimulated	 	 		pCMVSport
H0316	I/C	with poly I/C				3.0
H0521	Primary Dendritic Cells,	Primary Dendritic				pCMVSport
110521	lib 1	cells _		_		3.0
H0522	Primary Dendritic	Primary Dendritic				pCMVSport
110504	cells,frac 2	cells	ļ			3.0
H0524	Primary Dendritic Cells, CapFinder, frac 2	Primary Dendritic cells	1	{		pSport1
H0529	Myoloid Progenitor Cell	TF-1 Cell Line;				pCMVSport
	Line	Myoloid progenitor	1			3.0
		cell line				<u></u>
H0537	H. Primary Dendritic	Primary Dendritic				pCMVSport
	Cells,lib 3	cells				2.0
H0542	T Cell helper I	Helper T cell	{			pCMVSport
110542	T cell below II	Helper T cell	 			3.0 pCMVSport
H0543	T cell helper II	Helper I cell				3.0
H0545	Human endonietrial	Human endometrial	 			pCMVSport
	stromal cells-treated with	stromal cells-treated	ŀ	1		3.0
	progesterone	with proge				
H0551	Human Thymus Stromal	Human Thymus				pCMVSport
	Cells	Stromal Cells				3.0
H0552	Signal trap,Femur Bone	Femur Bone				Other
	Marrow,pooled	marrow, pooled from 8 male/female	ł	•		
H0556	Activated T-	T-Cells	Blood	Cell Line		Uni-ZAP
110330	cell(12h)/Thiouridine-re-	1-00113	Blood	Cen Eme		XR
H0560	excision KMH2	KMH2	 			pCMVSport
110300	1211112					3.0
H0575	Human Adult	Human Adult	Lung			Uni-ZAP
	Pulmonary;re-excision	Pulmonary				XR
H0576	Resting T-Cell; re-	· T-Cells	Blood	Cell Line		Lambda
	excision		ļ <u>_</u> _	 		ZAPII
H0578	Human Fetal Thymus	Fetal Thymus	Thymus			pSport1
H0580	Dendritic cells, pooled	Pooled dendritic				pCMVSport 3.0
H0581	Human Bone Marrow,	cells Human Bone	Bone Marrow			pCMVSport
10001	treated	Marrow	Done wantow			3.0
H0583	B Cell lymphoma	B Cell Lymphoma	B Cell		disease	pCMVSport
						3.0
H0584	Activated T-cells, 24	Activated T-Cells	Blood	Cell Line		Uni-ZAP
	hrs,re-excision	1	7.	<u> </u>	<u>-</u>	XR
H0585	Activated T-Cells,12	Activated T-Cells	Blood	Cell Line		Uni-ZAP XR
H0589	hrs,re-excision CD34 positive cells (cord	CD34 Positive Cells	Cord Blood			ZAP Express
בסנטנו	blood), re-ex	CD34 LOSITIVE CEITS	Cold Blood			LAN Express
H0591	Human T-cell	T-Cell Lymphoma	T-Cell		discase	Uni-ZAP
	lymphoma;re-excision					XR
H0606	Human Primary Breast	Human Primary	Breast		disease	Uni-ZAP
	Cancer;re-excision	Breast Cancer				XR
H0607	H.Leukocytes, normalized	H.Leukocytes				pCMVSport
TIOCOG	cot 50A3	LI Contraction				pCMVSport
H0608	H. Leukocytes, control	H.Leukocytes				l pcivi v sport
H0609	H. Leukocytes,	H.Leukocytes				pCMV Sport

	normalized cot > 500A	T				1
H0610	H. Leukocytes, normalized cot 5A	H.Leukocytes			!	pCMVSport
H0611	H. Leukocytes, normalized cot 500 B	H.Leukocytes				pCMVSport
H0612	H.Leukocytes, normalized cot 50 B	H.Leukocytes				pCMVSport
H0614	H. Leukocytes, normalized cot 500 A	H.Leukocytes				pCMVSport
H0625	Ku 812F Basophils Line	Ku 812F Basophils				pSport1
H0635	Human Activated T-Cells, re-excision.	Activated T-Cells	Blood	Cell Line		Uni-ZAP XR
H0637	Dendritic Cells From CD34 Cells	Dentritic cells from CD34 cells				pSport1
H0638	CD40 activated monocyte dendridic cells	CD40 activated monocyte dendridic cells				pSport1
H0641	LPS activated derived dendritic cells	LPS activated monocyte derived dendritic cells				pSport1
H0650	B-Cells	B-Cells				pCMVSport 3.0
H0653	Stromal Cells	Stromal Cells				pSport1
H0656	B-cells (unstimulated)	B-cells (unstimulated)				pSport1
H0657	B-cells (stimulated)	B-cells (stimulated)		ļ		pSport1
H0659	Ovary, Cancer (15395A1F): Grade II Papillary Carcinoma	Grade II Papillary Carcinoma, Ovary	Ovary		disease	pSport1
H0660	Ovary, Cancer: (15799A1F) Poorly differentiated carcinoma	Poorly differentiated carcinoma, ovary			disease	pSport1
H0673	Human Prostate Cancer, Stage B2; re-excision	Human Prostate Cancer, stage B2	Prostate			Uni-ZAP XR
H0677	TNFR degenerate oligo	B-Cells				PCRII
H0679	screened clones from Tonsil library	Human Tonsils				Other
\$0002	Мопосуtе activated	Monocyte-activated	blood	Cell Line	•	Uni-ZAP XR
\$0026	Stromal cell TF274	stromal cell	Bone marrow	Cell Line		Uni-ZAP XR
\$0027	Smooth muscle, serum treated	Smooth muscle	Pulmanary artery	Cell Line		Uni-ZAP XR
\$0031	Spinal cord	Spinal cord	spinal cord			Uni-ZAP XR
S0052	neutrophils control	human neutrophils	blood	Cell Line		Uni-ZAP XR
S0053	Neutrophils IL-1 and LPS induced	human neutrophil induced	blood	Cell Line		Uni-ZAP XR
S0114	Anergic T-cell	Anergic T-cell		Cell Line		Uni-ZAP XR
S0116	Bone marrow	Bone marrow	Bone marrow			Uni-ZAP XR
S0134	Apoptotic T-cell	apoptotic cells		Cell Line		Uni-ZAP XR
S0140	eosinophil-IL5 induced	eosinophil	lung	Cell Line		Uni-ZAP XR
S0142	Macrophage-oxLDL	macrophage- oxidized LDL treated	blood	Cell Line		Uni-ZAP XR
S0144	Macrophage (GM-CSF treated)	Macrophage (GM- CSF treated)				Uni-ZAP XR

S0180	Bone Marrow Stroma,	Bone Marrow			disease	Uni-ZAP
	TNF&LPS ind	Stroma, TNF & LPS induced				XR
S0182	Human B Cell 8866	Human B- Cell 8866				Uni-ZAP XR
S0196	Synovial IL-1/TNF stimulated	Synovial Fibroblasts				pSport1
S0212	Bone Marrow Stromal Cell, untreated	Bone Marrow Stromal Cell,untreated				pSport1
S0216	Neutrophils IL-1 and LPS induced	human neutrophil induced	blood	Cell Line		Uni-ZAP XR
S0218	Apoptotic T-cell, re- excision	apoptotic cells		Cell Line		Uni-ZAP XR
50222	H. Frontal cortex,epileptic;re- excision	H. Brain, Frontal Cortex, Epileptic	Brain		disease	Uni-ZAP XR
S0278	H Macrophage (GM-CSF treated), re-excision	Macrophage (GM- CSF treated)				Uni-ZAP XR
S0282	Brain Frontal Cortex, re- excision	Brain frontal cortex	Brain			Lambda ZAP II
S0292	Osteoarthritis (OA-4)	Human Osteoarthritic Cartilage	Bone		disease	pSport1
S0298	Bone marrow stroma, treated	Bone marrow stroma,treatedSB	Bone marrow			pSport1
S0308	Spleen/normal	Spleen normal				pSport1
S0314	Human osteoarthritis;fraction I	Human osteoarthritic cartilage			disease	pSport1
S0344	Macrophage-oxLDL; re- excision	macrophage- oxidized LDL treated	blood	Cell Line		Uni-ZAP XR
S0358	Colon Normal III	Colon Normal	Colon			pSport1
S0418	CHME Cell Line;treated 5 hrs	CHME Cell Line; treated				pCMVSport 3.0
S0426	Monocyte activated; re- excision	Monocyte-activated	blood	Cell Line		Uni-ŽAP XR
S0428	Neutrophils control; re- excision	human neutrophils	blood	Cell Line		Uni-ZAP XR
S0442	Colon Normal	Colon Normal				pSport1
S0452	Thymus	Thymus				pSport1
S0474	Human blood platelets	Platelets	Blood platelets			Other
S3016	Basophil	Basophil TH1 cells				Uni-ZAP XR Uni-ZAP
\$3018	TH1 cells					XR Uni-ZAP
\$3020	TH2 cells	TH2 cells Human Manic	Brain		disease	XR Uni-ZAP
S6028	Human Manic Depression Tissue Activated T-cells	depression tissue	Blood	Cell Line	uiscase	XR pBluescript
T0002		Activated T-Cell, PBL fraction	Dioon	Cen Line		SK-
T0041	Jurkat T-cell G1 phase	Jurkat T-cell				pBluescript SK-
T0042	Jurkat T-Cell, S phase	Jurkat T-Cell Line				pBluescript SK-
T0071	Human Bone Marrow	Human Bone Магтоw				pBluescript SK-
L0002	Atrium cDNA library Human heart					

		·				
L0004	ClonTech HL 1065a					
L0005	Clontech human aorta			1	i	1
	polyA+ mRNA (#6572)	l			į	
L0017	Human (J. Swensen)					
L0020	Human activated dendritic				 	
L0020	cell mRNA	ļ i		ŀ	1	İ
7.0001						
L0021	Human adult (K.Okubo)				<u> </u>	
L0036	Human chromosome			İ	ļ	
	21q22 mRNA					
L0040	Human colon mucosa					
L0041	Human epidermal					1
	keratinocyte					<u> </u>
L0051	Human mRNA (Tripodis					
	and Ragoussis)			ŀ		
L0055	Human promyelocyte			 		·
L0060	Human thymus NSTH II		·-·			·
						
L0065	Liver HepG2 cell line.			 		
L0070	Selected chromosome 21			ļ		
	cDNA library					
L0109	Human brain cDNA	brain			<u> </u>	<u> </u>
L0142	Human placenta cDNA	placenta			ļ	
	(TFujiwara)		·		<u></u>	<u></u>
L0143	Human placenta polyA+	placenta				
	(TFujiwara)	Langame.			!	
L0157	Human fetal brain		brain	- 		
LUIJ	(TFujiwara)		oran.	1		
L0163	Human heart cDNA		heart			
10103			licatt	ł	ĺ	
	(YNakamura)	<u> </u>		1540		
L0171	Human lung	lung		A549	ł	
	adenocarcinoma A549	adenocarcinoma		<u> </u>		
L0177	Human newborn			Clonetics	1	İ
	melanocytes (T.Vogt)	i I		Corp.		
				(San		·
		i		Diego,	f	
		[]		CA)	į.	
				strain #68	j	
				and 2486		
L0239	Homo sapiens brain fetus	brain				
L0307	Human C3-A11N			C3-		
10307	Hullian C3-ATTN			Alin;		
	·	[1
				clonally	İ	<u> </u>
				related		Į.
				variant of	ĺ	1
				OCI	1	
			· · · · · · · · · · · · · · · · · · ·	LY8-C3P	<u></u>	
L0309	Human E8CASS	breast		E8CASS;	1]
		adenocarcinoma		variant of	l	I
				MCF7_	<u></u> .	<u></u>
L0352	Normalized infant brain,					BA, M13-
	Bento Soares			1	1	derived
L0361	Stratagene ovary		ovary			Bluescript
FOOT	_		57413		}	SK
1.0262	(#937217)		·	-	 	Bluescript
L0362	Stratagene ovarian cancer			İ]	
	(#937219)			 	ļ	SK-
L0363	NCI_CGAP_GC2	germ cell tumor		ł		Bluescript
					<u> </u>	SK-
L0365	NCI_CGAP_Phe1	pheochromocytoma				Bluescript
		·			1	SK
L0366	Stratagene schizo brain	schizophrenic brain			<u> </u>	Bluescript
F0300		S-11 frontal lobe		İ	ŀ	SK-
102/2	S11	Schwannoma tumor		- 	 	Bluescript
L0367	NCI_CGAP_Sch1	Schwamhoina tumor		1	i	
		1		1	I	SK

T				
NCI_CGAP_SS1	synovial sarcoma			Bluescript SK-
NCI_CGAP_AA1	adrenal adenoma	adrenal gland		Bluescript SK-
Johnston frontal cortex	pooled frontal lobe	brain		Bluescript
NCl_CGAP_Kid6	kidney tumor	kidney		SK- Bluescript
NCI_CGAP_Larl	larynx	larynx		SK- Bluescript
	<u></u>			SK-
	squamous cell carcinoma from vocal cord	larynx		Bluescript SK-
NCI_CGAP_Lu1	lung tumor	lung		Bluescript SK-
NCI_CGAP_HN4	squamous cell carcinoma	pharynx		Bluescript SK-
NCI_CGAP_Pr24	invasive tumor (cell line)	prostate		Bluescript SK-
NCI_CGAP_Pr23	prostate tumor	prostate		Bluescript SK-
NCI_CGAP_Gas1	gastric tumor	stomach		Bluescript SK-
NCI_CGAP_HN3	squamous cell carcinoma from base of tongue	tongue		Bluescript SK-
NCI_CGAP_GCB0	germinal center B- cells	tonsil		Bluescript SK-
NCI_CGAP_HN5	normal gingiva (cell line from primary			Bluescript SK-
H, Human adult Brain Cortex tissue				gtll
				Lafmid BA
Infant brain, LLNL array				lafmid BA
normalized infant brain	total brain	brain		lafmid BA
		whole brain		Lafmid BA
N3HFLSK20				Lafmid K
Chromosome 21, K. Gardiner				Lambda
Human retina cDNA randomly primed sublibrary	retina	еус		lambda gt10
Human retina cDNA Tsp509I-cleaved sublibrary	retina	eye		lambda gt10
multi-tissue normalized short-fragment	multi-tissue	pooled		lambda gt10
Adult heart, Clontech				Lambda gt l i
WATMI				lambda gt1 l
TEST1, Human adult Testis tissue				lambda nm 1149
HE6W				lambda zap
Human fetal heart, Lambda ZAP Express				Lambda ZAP Express
KG1-a Lambda Zap			KG1-a	Lambda Zap
Express cDNA library	}			Express (Stratagene)
Fetal brain, Stratagene				Lambda
	Johnston frontal cortex NCI_CGAP_Kid6 NCI_CGAP_Lar1 NCI_CGAP_Lar1 NCI_CGAP_Lu1 NCI_CGAP_Lu1 NCI_CGAP_HN4 NCI_CGAP_Pr24 NCI_CGAP_Pr23 NCI_CGAP_Gas1 NCI_CGAP_Gas1 NCI_CGAP_GB0 NCI_CGAP_HN3 NCI_CGAP_HN5 H, Human adult Brain Cortex tissue b4HB3MA Cot8-HAP-Ft Infant brain, LLNL array of Dr. M. Soares 1NIB normalized infant brain cDNA Soares infant brain 1NIB N3HFLSK20 Chromosome 21, K. Gardiner Human retina cDNA randomly primed sublibrary Human retina cDNA Tsp509I-cleaved sublibrary Human retina cDNA Tsp509I-cleaved sublibrary Human retina cDNA Tsp509I-cleaved sublibrary Human retina cDNA Tsp509I-cleaved sublibrary Human retina cDNA Tsp509I-cleaved sublibrary Human retina cDNA Tsp509I-cleaved sublibrary Human fetal heart, Lambda Zap Express KG1-a Lambda Zap Express KG1-a Lambda Zap Express KG1-a Lambda Zap Express cDNA library	NCI_CGAP_AA1 adrenal adenoma	NCI_CGAP_AA1 adrenal adenoma adrenal gland	NCI_CGAP_AA1 adrenal adenoma adrenal gland Johnston frontal cortex pooled frontal lobe brain NCI_CGAP_Kid6 kidney tumor kidney NCI_CGAP_Larl larynx larynx NCI_CGAP_Larl larynx larynx NCI_CGAP_HN2 squamous cell carcinoma from NCI_CGAP_Lul lung tumor lung NCI_CGAP_HN4 squamous cell pharynx Carcinoma carcinoma NCI_CGAP_P724 invasive tumor (cell line) NCI_CGAP_P723 prostate tumor prostate NCI_CGAP_Gas1 gastric tumor stomach NCI_CGAP_HN3 squamous cell carcinoma from base of tongue NCI_CGAP_HN3 squamous cell carcinoma from base of tongue NCI_CGAP_GBO germinal center B cells NCI_CGAP_HN5 normal gingiva (cell line from primary keratinocyt H, Human adult Brain Cortex tissue HHB3MA Cot8-HAP-Ft Infant brain, LLNL array of Dr. M. Soares INIB normalized infant brain tolal brain cDNA Soares infant brain 1NIB N3HFLSK20 Chromosome 21, K. Gardiner Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary Human retina cDNA retina eye sublibrary

L0480	Stratagene cat#937212	T	1		Lambda
L0460	(1992)	İ]	ZAP.
l	(1992)		1		pBluescript
		1			SK(-)
L0481	CD34+DIRECTIONAL				Lambda
					ZAPII
L0483	Human pancreatic islet	<u> </u>			Lambda
L					ZAPII
L0485	STRATAGENE Human	skeletal muscle	leg muscle		Lambda
	skeletal muscle cDNA				ZAPII
	library, cat. #936215.	ļ			
L0492	Human Genomic	<u> </u>	<u> </u>	L	pAMP
L0493	NCI_CGAP_Ov26	papillary serous	ovary		pAMP1
		carcinoma		<u> </u>	
L0498	NCI_CGAP_HSC3	CD34+, T negative,	bone marrow		pAMPI
	1	patient with chronic	[
1.0506	NGV CCAP P 16	myelogenou			4) (D)
L0506	NCI_CGAP_Br16	lobullar carcinoma	breast		pAMP1
1.0500	NCI_CGAP_Lu25	in situ bronchioalveolar	100-	 	pAMPI
L0508	NCI_CGAP_Lu25	carcinoma	lung		PAMPI
L0509	NCI_CGAP_Lu26	invasive	lung		pAMP1
LU309	NCI_COAF_Luzo	adenocarcinoma	lung		PAME 1
L0512	NCI_CGAP_Ov36	borderline ovarian	ovary	 	pAMP1
203.2	1101_00711_0130	carcinoma) 014,		J P
L0513	NCI_CGAP_Ov37	early stage papillary	ovary		pAMP1
		serous carcinoma		1	
L0515	NCI_CGAP_Ov32	papillary serous	ovary		pAMP1
		carcinoma	,		•
L0517	NCI_CGAP_Pr1				pAMP10
L0518	NCI_CGAP_Pr2				pAMP10
L0519	NCI_CGAP_Pr3				pAMP10
L0520	NCI_CGAP_Alv1	alveolar			pAMP10
		rhabdomyosarcoma			
L0521	NCI_CGAP_Ew1	Ewing"s sarcoma		<u> </u>	pAMP10
L0523	NCI_CGAP_Lip2	liposarcoma			pAMP10
L0526	NCI_CGAP_Pr12	metastatic prostate			pAMP10
		bone lesion			
L0527	NCI_CGAP_Ov2	ovary			pAMP10
L0528	NCI_CGAP_Pr5	prostate			pAMP10
L0529	NCI_CGAP_Pr6	prostate			pAMP10
L0530	NCI_CGAP_Pr8	prostate		ļ	pAMP10
L0532	NCI_CGAP_Thy1	thyroid	1	ļ	pAMP10
L0534	Chromosome 7 Fetal	brain	brain	J •	pAMP10
L0535	Brain cDNA Library NCI CGAP_Br5	infiltrating ductal	breast		pAMP10
E0333	NCI_COAF_BIS	carcinoma	bicast		pairti to
L0539	Chromosome 7 Placental	Carcinoma	placenta	 	pAMP10
20333	cDNA Library		Praventa		P. M. I.V
L0541	NCI_CGAP_Pr7	low-grade prostatic	prostate	 	pAMP10
203 (1		neoplasia			
L0542	NCI_CGAP_Pr11	normal prostatic	prostate		pAMP10
-	-	epithelial cells			
L0543	NCI_CGAP_Pr9	normal prostatic	prostate		pAMP10
		epithelial cells			
L0545	NCI_CGAP_Pr4.1	prostatic	prostate		pAMP10
		intraepithelial			
		neoplasia - high		1	
		grade			
L0546	NCI_CGAP_Pr18	stroma	prostate		pAMP10
L0547	NICT CCAD Date	tumor	prostate	. !	pAMP10
L0551	NCI_CGAP_Pr16 NCI_CGAP_HN7	normal squamous	prostato		pAMP10

		epithelium, floor of			
10556	NOLOCAR L-241	mouth	1	 -	-41/010
L0556 L0562	NCI_CGAP_Lu34.1 Chromosome 7 HeLa	large cell carcinoma	lung	HeLa cell	pAMP10 pAMP10
LU362	cDNA Library			line;	PAMPIO
L0565	Normal Human Trabecular Bone Cells	Bone	Hip		pBluescript
L0581	Stratagene liver (#937224)		liver		pBluescript SK
L0586	HTCDLI				pBluescript SK(-)
L0588	Stratagene endothelial cell 937223				pBluescript SK-
L0589	Stratagene fetal retina 937202				pBluescript SK-
L0590	Stratagene fibroblast (#937212)				pBluescript SK-
L0591	Stratagene HeLa cell s3 937216				pBluescript SK-
L0592	Stratagene hNT neuron (#937233)				pBluescript SK-
L0593	Stratagene neuroepithelium (#937231)				pBluescript SK-
L0594	Stratagene neuroepithelium NT2RAMI 937234				pBluescript SK-
L0595	Stratagene NT2 neuronal precursor 937230	neuroepithelial cells	brain		pBluescript SK-
L0596	Stratagene colon (#937204)		colon		pBluescript SK-
L0597	Stratagene corneal stroma (#937222)		comea		pBluescript SK-
L0598	Morton Fetal Cochlea	cochlea	ear		pBluescript SK-
L0599	Stratagene lung (#937210)		lung		pBluescript SK-
L0600	Weizmann Olfactory Epithelium	olfactory epithelium	nose		pBluescript SK-
L0601	Stratagene pancreas (#937208)		pancreas		pBluescript SK-
L0602	Pancreatic Islet	pancreatic islet	pancreas	ļ	pBluescript SK-
L0603	Stratagene placenta (#937225)		placenta		pBluescript SK-
L0604	Stratagene muscle 937209	muscle	skeletal muscle		pBluescript SK-
L0605	Stratagene fetal spleen (#937205)	fetal spleen	spleen		pBluescript SK-
L0606	NCI_CGAP_Lym5	follicular lymphoma	lymph node		pBluescript SK-
L0607	NCI_CGAP_Lym6	mantle cell lymphoma	lymph node		pBluescript SK-
L0608	Stratagene lung carcinoma 937218	lung carcinoma	lung	NCI-H69	pBluescript SK-
L0612	Schiller oligodendroglioma	oligodendroglioma	brain		pBluescript SK- (Stratagene)
L0617	Chromosome 22 exon				pBluescript[[KS+
L0619	Chromosome 9 exon II				pBluescriptII

					KS+
L0622	HMI				pcDNAII
<u> </u>					(Invitrogen)
L0623	HM3	pectoral muscle]	pcDNAII
1000	NGI COAD CCI	(after mastectomy)			(Invitrogen)
L0626	NCI_CGAP_GC1	bulk germ cell seminoma	l	l	pCMV- SPORT2
L0629	NCI_CGAP_Mel3	metastatic	bowel (skin	 	pCMV-
20027	1101_00111	melanoma to bowel	primary)	1	SPORT4
L0631	NCI_CGAP_Br7		breast		pCMV-
			<u> </u>		SPORT4
L0633	NCI_CGAP_Lu6	small cell carcinoma	lung	1	pCMV-
10024	NCI_CGAP_Ov8				SPORT4
L0634	NCI_CGAP_OVO	serous adenocarcinoma	ovary	•	pCMV- SPORT4
L0635	NCI_CGAP_PNS1	dorsal root ganglion	peripheral	 	pCMV-
		3 3	nervous		SPORT4
			system		
L0637	NCI_CGAP_Brn53	three pooled	brain		pCMV-
10622	1101 001 0 0	meningiomas		<u> </u>	SPORT6
L0638	NCI_CGAP_Bm35	tumor, 5 pooled (see description)	brain		pCMV- SPORT6
L0639	NCI_CGAP_Bm52	tumor, 5 pooled (see	brain	 	pCMV-
1 20035	iver_com_bilist	description)) orani		SPORT6
L0640	NCI_CGAP_Br18	four pooled high-	breast		pCMV-
		grade tumors,			SPORT6
		including two prima	ļ		
L0641	NCI_CGAP_Co17	juvenile granulosa	colon		pCMV- SPORT6
L0643	NCI_CGAP_Co19	tumor moderately	colon		pCMV-
L0043	NCI_COAI_COIS	differentiated	Colon		SPORT6
	l	adenocarcinoma		l L	
L0644	NCI_CGAP_Co20	moderately	colon		pCMV-
		differentiated	ì		SPORT6
L0645	NCL CCAR Coll	adenocarcinoma	colon	 	pCMV-
1.0643	NCI_CGAP_Co21	moderately differentiated	Colon]]	SPORT6
		adenocarcinoma			
L0646	NCI_CGAP_Co14	moderately-	colon		pCMV-
		differentiated			SPORT6
	1	adenocarcinoma		 	- CD (IV
L0647	NCI_CGAP_Sar4	five pooled sarcomas, including	connective tissue]	pCMV- SPORT6
		myxoid liposarcoma	tissue		J. OK.
L0648	NCI_CGAP_Eso2	squamous cell	esophagus		pCMV-
		carcinoma			SPORT6
L0649	NCI_CGAP_GU1	2 pooled high-grade	genitourinary	[pCMV-
		transitional cell	tract		SPORT6
L0651	NCI_CGAP_Kid8	tumors renal cell tumor	kidney		pCMV-
ונטעב	INCI_COMI_IXIGO	tonai con tuittoi	Ridlicy]	SPORT6
L0652	NCI_CGAP_Lu27	four pooled poorly-	lung		pCMV-
		differentiated	_		SPORT6
		adenocarcinomas	<u> </u>	ļ	
L0653	NCI_CGAP_Lu28	two pooled	lung		pCMV- SPORT6
		squamous cell carcinomas			SPURIO
L0654	NCI_CGAP_Lu31	Carcinomas	lung, cell line		pCMV-
20034			, , , , , , , , , , , , , , , ,		SPORT6
L0655	NCI_CGAP_Lym12	lymphoma,	lymph node		pCMV-
		follicular mixed			SPORT6
		small and large cell			-0.97
L0656	NCI_CGAP_Ov38	normal epithelium	ovary	<u> </u>	pCMV-

		T		r	SDODT4
L0657	NCI_CGAP_Ov23	tumor, 5 pooled (see	ovary	 	SPORT6 pCMV-
		description)			SPORT6
L0658	NCI_CGAP_Ov35	tumor, 5 pooled (see description)	очагу		pCMV- SPORT6
L0659	NCI_CGAP_Pan1	adenocarcinoma	pancreas		pCMV- SPORT6
L0661	NCI_CGAP_Mel15	malignant melanoma, metastatic to lymph node	skin		pCMV- SPORT6
L0662	NCI_CGAP_Gas4	poorly differentiated adenocarcinoma with signet r	stomach		pCMV- SPORT6
L0663	NCI_CGAP_Ut2	moderately- differentiated endometrial adenocarcino	uterus		pCMV- SPORT6
L0664	NCI_CGAP_Ut3	poorly-differentiated endometrial adenocarcinoma,	uterus		pCMV- SPORT6
L0665	NCI_CGAP_Ut4	serous papillary carcinoma, high grade, 2 pooled t	uterus		pCMV- SPORT6
L0666	NCI_CGAP_Ut1	well-differentiated endometrial adenocarcinoma, 7	uterus		pCMV- SPORT6
L0667	NCI_CGAP_CML1	myeloid cells, 18 pooled CML cases, BCR/ABL rearra	whole blood		pCMV- SPORT6
L0682	Stanley Frontal NB pool 2	frontal lobe (see description)	brain		pCR2.1- TOPO (Invitrogen)
L0686	Stanley Frontal SN pool 2	frontal lobe (see description),	brain		pCR2.1- TOPO (Invitrogen)
L0697	Testis 1				PGEM 5zf(+)
L0700	Outward Alu-primed hncDNA library				pGEM-3Z
L0717	Gessler Wilms tumor				pSPORT1
L0720	PN001-Normal Human Prostate		prostate		pSportI
L0731	Soares_pregnant_uterus_ NbHPU		uterus		pT7T3-Pac
L0738	Human colorectal cancer				pT7T3D
L0740	Soares melanocyte 2NbHM	melanocyte			pT7F3D (Pharmacia) with a modified polylinker
L0741	Soares adult brain N2b4HB55Y		brain	·	pT7T3D (Pharmacia) with a modified polylinker
L0742	Soares adult brain N2b5HB55Y	·	brain		pT7T3D (Pharmacia) with a modified polylinker
L0743	Soares breast 2NbHBst		breast		pT7T3D (Pharmacia)

Γ	1		1		with a
					modified
	į.		1		polylinker
L0744	Soares breast 3NbHBst		breast		pT7T3D
			1		(Pharmacia)
	į				with a
			Ì]	modified
					polylinker
L0745	Soares retina N2b4HR	retina	eye		pT7T3D
			1		(Pharmacia)
				f	with a
		•		i I	modified
					polylinker
L0746	Soares retina N2b5HR	retina	eye		pT7T3D
					(Pharmacia) with a
	i				modified
		•	1		polylinker
L0747	Soares_fetal_heart_NbHH		heart	 	· pT7F3D
D0747	19W		, neare	[(Pharmacia)
	1211				with a
	ł		1		modified
		•			polylinker
L0748	Soares fetal liver spleen		Liver and		рТ7Г3D
	INFLS		Spleen		(Pharmacia)
				[with a
			Ì	i i	modified
				<u> </u>	polylinker
L0749	Soares_fetal_liver_spleen		Liver and		pT7T3D
	_1NFLS_S1		Spleen		(Pharmacia) with a
					modified
					polylinker
L0750	Soares_fetal_lung_NbHL1		lung		pT7T3D
L0730	9W		1 ''''		(Pharmacia)
			J		with a
					modified
					polylinker
L0751	Soares ovary tumor	ovarian tumor	ovary		pT7T3D
	NbHOT		Į.	ļ ļ	(Pharmacia)
					with a
				1	modified
		11		 	polylinker
L0752	Soares_parathyroid_tumor	parathyroid tumor	parathyroid gland		pT7T3D (Pharmacia)
	_NbHPA		granu		with a
					modified
			Ì		polylinker
L0753	Soares_pineal_gland_N3H		pineal gland		pT7T3D
	PG		' "		(Pharmacia)
					with a
			j		modified
					polylinker
L0754	Soares placenta Nb2HP		placenta		pT7T3D
					(Pharmacia)
					with a
			Ì		modified
. 0			1	 	polylinker -T7T2D
L0755	Soares_placenta_8to9wcc		placenta		pT7T3D (Pharmacia)
	ks_2NbHP8to9W		-		with a
			1		modified
					polylinker
	<u> </u>		<u> </u>	L	

	Ta					TTT 10
L0756	Soares_multiple_sclerosis	multiple sclerosis		i		pT7T3D
	_2NbHMSP	lesions				(Pharmacia)
						with a
• •					Ì	modified
1	İ			1		polylinker
				ļ		V_TYPE
L0757	Soares_senescent_fibrobla	senescent fibroblast		ļ		pT7T3D
	sts_NbHSF			ļ		(Pharmacia)
	i			1		with a
	1				İ	modified
İ						polylinker
				ŀ		V_TYPE
L0758	Soares_testis_NHT			i		pT7T3D-Pac
				İ	i	(Pharmacia)
ļ		1		1		with a
				1	Ĭ	modified
					1	polylinker
L0759	Soares_total_fetus_Nb2H	<u> </u>			i — —	pT7T3D-Pac
20137	F8_9w			ł	}	(Pharmacia)
	1.0_>			1	İ	with a
	1			ļ		modified
]	1			ł		polylinker
L0761	NCI_CGAP_CLL1	B-cell, chronic		 		pT7T3D-Pac
10/01	INCI_COMF_CELEI	lymphotic leukemia		Į.	1	(Pharmacia)
1		Tymphone reakemia			1	with a
Ì				{		modified
1				t		polylinker
1.07/2	NGL GCAR R-13	L				pT7T3D-Pac
L0762	NCI_CGAP_Br1.1	breast		j	i	(Pharmacia)
				1		with a
						modified
					•	
				 		polylinker
L0763	NCI_CGAP_Br2	breast		İ		pT7T3D-Pac
				ļ	ł	(Pharmacia)
				ł	Į.	with a
				1		modified
				 	ļ	polylinker
L0764	NCI_CGAP_Co3	colon				pT7T3D-Pac
ļ .					<u> </u>	(Pharmacia)
						with a
1						modified
						polylinker
L0766	NCI_CGAP_GCB1	germinal center B			1	рТ7Г3Д-Рас
		cell			1	(Pharmacia)
					1	with a
	İ					modified
						polylinker
L0767	NCI_CGAP_GC3	pooled germ cell		}		pT7T3D-Pac
		tumors		}	ł	(Pharmacia)
,				1	1	with a
				J	1	modified
				<u> </u>	<u> </u>	polylinker
L0768	NCI_CGAP_GC4	pooled germ cell				pT7F3D-Pac
		tumors			1	(Pharmacia)
]	with a
				-	1	modified
					ĺ	polylinker
L0769	NCI_CGAP_Brn25	anaplastic	brain	 	<u> </u>	pT7T3D-Pac
LU/09	INCI_CGMP_DIII23	oligodendroglioma	Viaili			(Pharmacia)
		. outhorising Suoma				with a
					1	modified
				1	1	polylinker
	1101 00 10 5		t •	 	 	
L0770	NCI_CGAP_Bm23	glioblastoma	brain			pT7T3D-Pac
i i		(pooled)		<u> </u>	L	(Pharmacia)

					
1	1		ł		with a
					modified
	<u> </u>				polylinker
L0771	NCI_CGAP_Co8	adenocarcinoma	colon		pT7T3D-Pac
ľ					(Pharmacia)
1				1	with a
İ	İ	1			modified
1				!	polylinker
L0772	NCI_CGAP_Co10	colon tumor RER+	colon		pT7T3D-Pac
					(Pharmacia)
1			1] !	with a
į.	1	}	}]	modified
1			}		polylinker
L0773	NCI_CGAP_Co9	colon tumor RER+	colon	 	pT7F3D-Pac
LUTTS	NCI_COAI_COS	COION MINO KERT	Colon	}	(Pharmacia)
ł					with a
			1] '	modified
	1				
10774	NOT CC 4 S IC IS	 		 	polylinker
L0774	NCI_CGAP_Kid3		kidney	1	pT7T3D-Pac
1	1			1	(Pharmacia)
	1		ł		with a
1	!		1	1 1	modified
		<u> </u>	l	L	polylinker
L0775	NCI_CGAP_Kid5	2 pooled tumors	kidney		pT7T3D-Pac
l		(clear cell type)			(Pharmacia)
1	ſ		ĺ	1	with a
1					modified
l	<u> </u>				polylinker
L0776	NCI_CGAP_Lu5	carcinoid	lung		pT7T3D-Pac
į			İ		(Pharmacia)
1	1		}	j i	with a
i		1	Ì		modified
					polylinker
L0777	Soares_NhHMPu_S1	Pooled human	mixed (see	 	pT7T3D-Pac
]		melanocyte, fetal	below)		(Pharmacia)
	İ	heart, and pregnant	""		with a
1	i	Industry unite programme			modified
]	i		polylinker
L0779	Soares_NFL_T_GBC_S1	· · · · · · · · · · · · · · · · · · ·	pooled	 	pT7T3D-Pac
1 20113	Boales_141 E_1_GDC_61		pooled	1	(Pharmacia)
1				1	with a
İ			1		modified
		{			polylinker
1.0790	C NCT FO OW OT		pooled	 	pT7T3D-Pac
L0780	Soares_NSF_F8_9W_OT		pooled	1	(Pharmacia)
	_PA_P_SI		l		with a
1			1]	modified
		1	1		
	No. 60 to 5 5 5			 	polylinker
L0783	NCI_CGAP_Pr22	normal prostate	prostate	1	pT7F3D-Pac
	1	}	1	1	(Pharmacia)
					with a
		1	1		modified
		ļ <u></u>			polylinker
L0784	NCI_CGAP_Lei2	leiomyosarcoma	soft tissue	1	pT7T3D-Pac
					(Pharmacia)
		[[1	with a
				1	modified
				1	polylinker
L0785	Barstead spleen HPLRB2		spleen		pT7f3D-Pac
			•		(Pharmacia)
					with a
	•				modified
				1	polylinker
		<u></u>	L		T borymines

	T =	γ	r			77770
L0786	Soares_NbHFB		whole brain	ļ		pT713D-Pac
						(Pharmacia) with a
				1		modified
						polylinker
10505	VOLCOAD S. L.	 				pT7T3D-Pac
L0787	NCI_CGAP_Sub1	<u> </u>]		(Pharmacia)
1		1		į		with a
				ì	ļ	modified
	Į			1		polylinker
L0788	NCI_CGAP_Sub2	 				pT7T3D-Pac
1 20,000	Mol_com_bus					(Pharmacia)
						with a
			ŀ	ļ		modified
				<u></u>		polylinker
L0789	NCI_CGAP_Sub3			1	1	pT7T3D-Pac
					1	(Pharmacia)
					•	with a
						modified
		<u> </u>				polylinker
L0790	NCI_CGAP_Sub4	İ		Ì		pT7T3D-Pac
]		(Pharmacia) with a
				ł		modified
		1	ļ	1		polylinker
1.0701	NCI_CGAP_Sub5					pT7T3D-Pac
L0791	NCI_CGAP_Sub3		ļ	Ì		(Pharmacia)
			,			with a
			Ì	1		modified
						polylinker
L0792	NCI_CGAP_Sub6					pT7T3D-Pac
				ļ	l	(Pharmacia)
	}		1	i	Ì	with a
						modified
						polylinker
L0794	NCI_CGAP_GC6	pooled germ cell				pT7T3D-Pac
		tumors			ļ	(Pharmacia) with a
					1	modified
		•				polylinker
L0796	NCI_CGAP_Bm50	medulloblastoma	brain	 	 	pT7T3D-Pac
10/30	NCI_COAF_BIIDO	Incumoolasiona	"""]	(Pharmacia)
				İ		with a
<u> </u>				l		modified
						polylinker
L0800	NCI_CGAP_Co16	colon tumor, RER+	colon			pT7T3D-Pac
	<u>-</u>			İ		(Pharmacia)
						with a
			ŀ	1		modified
		ļ			 	polylinker
L0803	NCI_CGAP_Kid11		kidney			pT7T3D-Pac
						(Pharmacia) with a
						modified
1		1	[polylinker
1.0004	NCL CGAD VIAIS	2 pooled tumors	kidney			pT7T3D-Pac
L0804	NCI_CGAP_Kid12	(clear cell type)	Mulicy			(Pharmacia)
}		(crear cen type)	l	1		with a
			ļ	}		modified
1						polylinker
L0805	NCI_CGAP_Lu24	carcinoid	lung			pT7T3D-Pac
	1			Į.	}	(Pharmacia)
		İ	l			with a
						modified

[1	polylinker
L0806	NCI_CGAP_Lu19	squamous cell carcinoma, poorly differentiated (4	lung	pT7T3D-Pac (Pharmacia) with a modified polylinker
L0809	NCI_CGAP_Pr28		prostate	pT7T3D-Pac (Pharmacia) with a modified polylinker
L2245	NEM subtracted human fetal kidney cDNA			pUEX1
1.2250	Human cerebral cortex	cerebral cortex		
L2251	Human fetal lung	Fetal lung		

TABLE 5

OMIM	Description
Reference	
100678	ACAT2 deficiency
102578	Leukemia, acute promyelocytic, PML/RARA type
102770	Myoadenylate deaminase deficiency
103000	Hemolytic anemia due to adenylate kinase deficiency
103050	Autism, succinylpurinemic
103050	Adenylosuccinase deficiency
103600	[Dysalbuminemic hyperthyroxinemia]
103600	[Dysalbuminemic hyperzincemia], 194470
103600	Analbuminemia
103850	Aldolase A deficiency
104150	[AFP deficiency, congenital]
104150	[Hereditary persistence of alpha-fetoprotein]
104170	NAGA deficiency, mild
104170	Schindler disease
104170	Kanzaki disease
104311	Alzheimer disease-3
104500	Amelogenesis imperfecta-2, hypoplastic local type
104770	Amyloidosis, secondary, susceptibility to
105580	Anal canal carcinoma
106165	Hypertension, essential, 145500
106300	Ankylosing spondylitis
107280	Cerebrovascular disease, occlusive
107280	Alpha-1-antichymotrypsin deficiency
107300	Antithrombin III deficiency
107400	Emphysema
107400	Emphysema-cirrhosis
107470	Atypical mycobacterial infection, familial disseminated, 209950
107470	BCG infection, generalized familial
107470	Tuberculosis, susceptibility to
107670	Apolipoprotein A-II deficiency
107741	Hyperlipoproteinemia, type III
107777	Diabetes insipidus, nephrogenic, autosomal recessive, 222000
108725	Atherosclerosis, susceptibility to
108730	Brody myopathy, 601003
108800	Atrial septal defect, secundum type
108962	Hypertension, salt-resistant
108985	Atrophia areata
109150	Machado-Joseph disease
109270	Renal tubular acidosis, distal, 179800
109270	Spherocytosis, hereditary
109270	[Acanthocytosis, one form]
109270	[Elliptocytosis, Malaysian-Melanesian type]

109270	Hemolytic anemia due to band 3 defect
109560	Leukemia/lymphoma, B-cell, 3
109690	Asthma, nocturnal, susceptibility to
109690	Obesity, susceptibility to
109700	Hemodialysis-related amyloidosis
110700	Vivax malaria, susceptibility to
113705	Ovarian cancer
113705	Breast cancer-1
113721	Breast cancer
113900	Heart block, progressive familial, type I
114208	Malignant hyperthermia susceptibility 5, 601887
114208	Hypokalemic periodic paralysis, 170400
114350	
114835	Leukemia, acute myeloid
	Monocyte carboxyesterase deficiency
115470	Cat eye syndrome
116800	Cataract, Marner type
116860	Cavernous angiomatous malformations
117700	[Hypoceruloplasminemia, hereditary]
117700	Hemosiderosis, systemic, due to aceruloplasminemia
118210	Charcot-Marie-Tooth neuropathy-2A
118485	Polycystic ovary syndrome with hyperandrogenemia
119300	van der Woude syndrome
120110	Metaphyseal chondrodysplasia, Schmid type
120215	Ehlers-Danlos syndrome, type I, 130000
120215	Ehlers-Danlos syndrome, type II, 130010
120220	Bethlem myopathy, 158810
120240	Bethlem myopathy, 158810
120260	Epiphyseal dysplasia, multiple, type 2, 600204
120280	Stickler syndrome, type III
120280	Marshall syndrome, 154780
120290	OSMED syndrome, 215150
120290	Stickler syndrome, type II, 184840
120435	Muir-Torre syndrome, 158320
120435	Colorectal cancer, hereditary, nonpolyposis, type 1 Ovarian
	cancer
120550	Clq deficiency, type A
120570	Clq deficiency, type B
120575	Clq deficiency, type C
120620	SLE susceptibility
120620	CR1 deficiency
120700	C3 deficiency
120810	C4 deficiency
120820	C4 deficiency
120900	C5 deficiency
120920	Measles, susceptibility to
120940	C9 deficiency
	

120070	
120950	C8 deficiency, type I
120960	C8 deficiency, type II
121011	Deafness, autosomal dominant 3, 601544
121011	Deafness, autosomal recessive 1, 220290
121014	Heterotaxia, visceroatrial, autosomal recessive
121360	Myeloid leukemia, acute, M4Eo subtype
121700	Congenital hereditary endothelial dystrophy of cornea
121800	Corneal dystrophy, crystalline, Schnyder
122000	Corneal dystrophy, posterior polymorphous
122500	[Transcortin deficiency]
122560	ACTH deficiency, 201400
122720	Nicotine addiction, protection from
122720	Coumarin resistance, 122700
123000	Craniometaphyseal dysplasia
123101	Craniosynostosis, type 2
123270	[Creatine kinase, brain type, ectopic expression of]
123580	Cataract, congenital, autosomal dominant
123620	Cataract, cerulean, type 2, 601547
123829	Melanoma
123940	White sponge nevus, 193900
124030	Parkinsonism, susceptibility to
124030	Debrisoquine sensitivity
124200	Darier disease (keratosis follicularis)
125490	Dentinogenesis imperfecta-1
125852	Insulin-dependent diabetes mellitus-2
126060	Anemia, megaloblastic, due to DHFR deficiency
126340	Xeroderma pigmentosum, group D, 278730
126391	DNA ligase I deficiency
126452	Autonomic nervous system dysfunction
126452	[Novelty seeking personality]
126600	Drusen, radial, autosomal dominant
126650	Chloride diarrhea, congenital, Finnish type, 214700
126650	Colon cancer
129490	Ectodermal dysplasia-3, anhidrotic
129500	Ectodermal dysplasia, hidrotic
129900	EEC syndrome-1
130410	Glutaricaciduria, type IIB
130500	Elliptocytosis-1
130650	Beckwith-Wiedemann syndrome
131195	Hereditary hemorrhagic telangiectasia-1, 187300
131210	Atherosclerosis, susceptibility to
131400	Eosinophilia, familial
132700	Cylindromatosis
133170	Erythremia
133171	[Erythrocytosis, familial], 133100
133200	Erythrokeratodermia variabilis
	

133780	Vitreoretinopathy, exudative, familial
134370	Factor H deficiency
134370	Hemolytic-uremic syndrome, 235400
134370	Membroproliferative glomerulonephritis
134570	Factor XIIIA deficiency
134580	Factor XIIIB deficiency
134790	Hyperferritinemia-cataract syndrome, 600886
135300	Fibromatosis, gingival
135940	Ichthyosis vulgaris, 146700
136132	[Fish-odor syndrome], 602079
136435	Ovarian dysgenesis, hypergonadotropic, with normal karyotype,
	233300
136550	Macular dystrophy, North Carolina type
136836	Fucosyltransferase-6 deficiency
137600	Iridogoniodysgenesis syndrome
138079	Hyperinsulinism, familial, 602485
138079	MODY, type 2, 125851
138140	Glucose transport defect, blood-brain barrier
138190	Diabetes mellitus, noninsulin-dependent
138491	Startle disease, autosomal recessive
138491	Startle disease/hyperekplexia, autosomal dominant, 149400
138491	Hyperekplexia and spastic paraparesis
138570	Non-insulin dependent diabetes mellitus, susceptibility to
138971	Kostmann neutropenia, 202700
138981	Pulmonary alveolar proteinosis, 265120
139190	Gigantism due to GHRF hypersecretion
139190	Isolated growth hormone deficiency due to defect in GHRF
139191	Growth hormone deficient dwarfism
139330	Night blindness, congenital stationary
139350	Epidermolytic hyperkeratosis, 113800
139350	Keratoderma, palmoplantar, nonepidermolytic
139360	Pituitary ACTH-secreting adenoma
140100	[Anhaptoglobinemia]
140100	[Hypohaptogloginemia]
141900	Methemoglobinemias, beta-
141900	Sickle cell anemia
141900	Thalassemias, beta-
141900	Erythremias, beta-
141900	HPFH, deletion type
141900	Heinz body anemias, beta-
142000	Thalassemia due to Hb Lepore
142000	Thalassemia, delta-
142200	HPFH, nondeletion type A
142250	HPFH, nondeletion type G
142270	Hereditary persistence of fetal hemoglobin
142360	Thrombophilia due to heparin cofactor II deficiency
176700	I imonioopinita dae to nepariti ootaotor ii detiotelley

142470	[Hereditary persistence of fetal hemoglobin, heterocellular]
142857	Pemphigoid, susceptibility to
142858	Beryllium disease, chronic, susceptibility to
142959	Hand-foot-uterus syndrome, 140000
143200	Wagner syndrome
143200	Erosive vitreoretinopathy
143890	Hypercholesterolemia, familial
144200	Epidermolytic palmoplantar keratoderma
145001	Hyperparathyroidism-jaw tumor syndrome
145260	
	Pseudohypoaldosteronism, type II
145981	Hypocalciuric hypercalcemia, type II
146760	[IgG receptor I, phagocytic, familial deficiency of]
146790	Lupus nephritis, susceptibility to
147061	Allergy and asthma susceptibility
147141	Leukemia, acute lymphoblastic
147280	Hepatocellular carcinoma
147440	Growth retardation with deafness and mental retardation
147557	Epidermolysis bullosa, junctional, with pyloric atresia, 226730
147570	Interferon, immune, deficiency
147575	Myelodysplastic syndrome, preleukemic .
147575	Myelogenous leukemia, acute
147575	Macrocytic anemia refractory, of 5q- syndrome, 153550
147781	Atopy, susceptibility to
148040	Epidermolysis bullosa simplex, Koebner, Dowling-Meara, and Weber-Cockayne types, 131900, 131760, 131800
148041	Pachyonychia congenita, Jadassohn-Lewandowsky type, 167200
148043	Meesmann corneal dystrophy, 122100
148065	White sponge nevus, 193900
148066	Epidermolysis bullosa simplex, Koebner, Dowling-Meara, and Weber-Cockayne types, 131900, 131760, 131800
148066	Epidermolysis bullosa simplex, recessive, 601001
148067	Nonepidermolytic palmoplantar keratoderma, 600962
148067	Pachyonychia congenita, Jadassohn-Lewandowsky type, 167200
148069	Pachyonychia congenita, Jackson-Lawler type, 167210
148070	Liver disease, susceptibility to, from hepatotoxins or viruses
148080	Epidermolytic hyperkeratosis, 113800
148900	Klippel-Feil syndrome with laryngeal malformation
150000	Exertional myoglobinuria due to deficiency of LDH-A
150210	Lactoferrin-deficient neutrophils, 245480
150240	Cutis laxa, marfanoid neonatal type
150250	Larsen syndrome, autosomal dominant
150270	Laryngeal adductor paralysis
150292	Epidermolysis bullosa, Herlitz junctional type, 226700
150310	Epidermolysis bullosa, Herlitz junctional type, 226700
150310	Epidermolysis bullosa, generalized atrophic benign, 226650
151385	Leukemia, acute myeloid
131303	Leuxenna, acute mycroru

151410	Leukemia, chronic myeloid
151440	Leukemia, T-cell acute lymphoblastoid
151670	Hepatic lipase deficiency
152200	Coronary artery disease, susceptibility to
152445	Vohwinkel syndrome, 124500
152445	Erythrokeratoderma, progressive symmetric, 602036
152760	Hypogonadotropic hypogonadism due to GNRH deficiency, 227200
152780	Hypogonadism, hypergonadotropic
152780	Male pseudohermaphroditism due to defective LH
152790	Precocious puberty, male, 176410
152790	Leydig cell hypoplasia
153455	Cutis laxa, recessive, type I, 219100
153880	
	Macular dystrophy, dominant cystoid
154275	Malignant hyperthermia susceptibility 2
154276	Malignant hyperthermia susceptibility 3
154500	Treacher Collins mandibulofacial dysostosis
154550	Carbohydrate-deficient glycoprotein syndrome, type Ib, 602579
156225	Muscular dystrophy, congenital merosin-deficient
157170	Holoprosencephaly-2
157640	PEO with mitochondrial DNA deletions, type 1
159000	Muscular dystrophy, limb-girdle, type 1A
159001	Muscular dystrophy, limb-girdle, type 1B
159440	Charcot-Marie-Tooth neuropathy-1B, 118200
159440	Dejerine-Sottas disease, myelin P-related, 145900
159440	Hypomyelination, congenital
160781	Cardiomyopathy, hypertrophic, mid-left ventricular chamber type
160900	Myotonic dystrophy
162200	Neurofibromatosis, type 1
162200	Watson syndrome, 193520
164040	Leukemia, acute promyelocytic, NPM/RARA type
164200	Oculodentodigital dysplasia
164200	Syndactyly, type III, 186100
164500	Spinocerebellar ataxia-7
164731	Ovarian carcinoma, 167000
164860	Renal cell carcinoma, papillary, familial and sporadic
164953	Liposarcoma
165215	3q21q26 syndrome
165500	Optic atrophy 1
166600	Osteopetrosis, AD, type II
166800	Otosclerosis
167000	Ovarian cancer, serous
167250	Paget disease of bone
167409	Optic nerve coloboma with renal disease, 120330
10/712	
167415	Hypothyroidism, congenital, due to thyroid dysgenesis or hypoplasia

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168360	Paraneoplastic sensory neuropathy
168468	Metaphyseal chondrodysplasia, Murk Jansen type, 156400
168610	Parkinsonism-dementia with pallidopontonigral degeneration
169600	Hailey-Hailey disease
170261	Bare lymphocyte syndrome, type I, due to TAP2 deficiency
170650	Periodontitis, juvenile
170995	Zellweger syndrome-2
171190	Hypertension, essential, 145500
171760	Hypophosphatasia, adult, 146300
171760	Hypophosphatasia, infantile, 241500
171860	Hemolytic anemia due to phosphofructokinase deficiency
172400	Hemolytic anemia due to glucosephosphate isomerase deficiency
172400	Hydrops fetalis, one form
172411	Colorectal cancer, resistance to
172471	Glycogenosis, hepatic, autosomal
172490	Phosphorylase kinase deficiency of liver and muscle, 261750
173350	Plasminogen Tochigi disease
173350	Plasminogen deficiency, types I and II
173350	Thrombophilia, dysplasminogenemic
173360	Thrombophilia due to excessive plasminogen activator inhibitor
173360	Hemorrhagic diathesis due to PAI1 deficiency
173610	Platelet alpha/delta storage pool deficiency
173850	Polio, susceptibility to
174000	Medullary cystic kidney disease, AD
174900	Polyposis, juvenile intestinal
176000	Porphyria, acute intermittent
176100	Porphyria cutanea tarda
176100	Porphyria, hepatoerythropoietic
176705	Breast cancer, sporadic
176730	Diabetes mellitus, rare form
176730	Hyperproinsulinemia, familial
176730	MODY, one form
176830	Obesity, adrenal insufficiency, and red hair
176830	ACTH deficiency
176860	Purpura fulminans, neonatal
176860	Thrombophilia due to protein C deficiency
176943	Apert syndrome, 101200
176943	Pfeiffer syndrome, 101600
176943	Beare-Stevenson cutis gyrata syndrome, 123790
176943	Crouzon craniofacial dysostosis, 123500
176943	Jackson-Weiss syndrome, 123150
176947	Selective T-cell defect
177070	Spherocytosis, hereditary, Japanese type
177070	Hermansky-Pudlak syndrome, 203300
177900	Psoriasis susceptibility-1
178300	Ptosis, hereditary congenital, 1

178640	Pulmonary alveolar proteinosis, congenital, 265120
179450	Ragweed sensitivity
179755	
	Renal cell carcinoma, papillary, 1
179820	[Hyperproreninemia]
180020	Retinal cone dystrophy-1
180071	Retinitis pigmentosa, autosomal recessive
180100	Retinitis pigmentosa-1
180104	Retinitis pigmentosa-9
180105	Retinitis pigmentosa-10
180250	Retinol binding protein, deficiency of
180297	Anemia, hemolytic, Rh-null, suppressor type, 268150
180380	Night blindness, congenital stationery, rhodopsin-related
180380	Retinitis pigmentosa, autosomal recessive
180380	Retinitis pigmentosa-4, autosomal dominant
180901	Malignant hyperthermia susceptibility 1, 145600
180901	Central core disease, 117000
181405	Scapuloperoneal spinal muscular atrophy, New England type
181430	Scapuloperoneal syndrome, myopathic type
181460	Schistosoma mansoni, susceptibility/resistance to
181510	Schizophrenia
182138	Anxiety-related personality traits
182280	Small-cell cancer of lung
182380	Glucose/galactose malabsorption
182381	Renal glucosuria, 253100
182500	Cataract, congenital
182600	Spastic paraplegia-3A
182601	Spastic paraplegia-4
182860	Pyropoikilocytosis
182860	Spherocytosis, recessive
182860	Elliptocytosis-2
185000	Stomatocytosis I
185430	Atherosclerosis, susceptibility to
185470	Myopathy due to succinate dehydrogenase deficiency
185800	Symphalangism, proximal
186580	Arthrocutaneouveal granulomatosis
186770	Leukemia, T-cell acute lymphocytic
186780	CD3, zeta chain, deficiency
186880	Leukemia/lymphoma, T-cell
186921	Leukemia, T-cell acute lymphoblastic
186960	Leukemia/lymphoma, T-cell
187040	Leukemia-1, T-cell acute lymphoblastic
188040	Thrombophilia due to thrombomodulin defect
188070	Bleeding disorder due to defective thromboxane A2 receptor
188400	Velocardiofacial syndrome, 192430
188400	DiGeorge syndrome
188826	Sorsby fundus dystrophy, 136900
100020	Language mattachity, 120200

189800	Propolamnoja/galamnoja
<u> </u>	Preeclampsia/eclampsia
189980	Leukemia, chronic myeloid
190000	Atransferrinemia
190020	Bladder cancer, 109800
190040	Dermatofibrosarcoma protuberans
190040	Giant-cell fibroblastoma
190040	Meningioma, SIS-related
190100	Geniospasm
190195	Ichthyosiform erythroderma, congenital, 242100
190195	Ichthyosis, lamellar, autosomal recessive, 242300
190198	Leukemia, T-cell acute lymphoblastic
190685	Down syndrome
190900	Colorblindness, tritan
191030	Nemaline myopathy-1, 161800
191044	Cardiomyopathy, familial hypertrophic
191045	Cardiomyopathy, familial hypertrophic, 2, 115195
191290	Segawa syndrome, recessive
191315	Insensitivity to pain, congenital, with anhidrosis, 256800
192090	Ovarian carcinoma
192090	Breast cancer, lobular
192090	Endometrial carcinoma
192090	Gastric cancer, familial, 137215
192500	Jervell and Lange-Nielsen syndrome, 220400
192500	Long QT syndrome-1
194071	Wilms tumor, type 2
194071	Adrenocortical carcinoma, hereditary, 202300
200150	Choreoacanthocytosis
200350	Acetyl-CoA carboxylase deficiency
201470	Acyl-CoA dehydrogenase, short-chain, deficiency of
201910	Adrenal hyperplasia, congenital, due to 21-hydroxylase
	deficiency
202200	Glucocorticoid deficiency, due to ACTH unresponsiveness
203310	Ocular albinism, autosomal recessive
203500	Alkaptonuria
204500	Ceroid-lipofuscinosis, neuronal 2, classic late infantile
205900	Anemia, Diamond-Blackfan
207750	Hyperlipoproteinemia, type Ib
207800	Argininemia
208100	Arthrogryposis multiplex congenita, neurogenic
208250	Jacobs syndrome
210900	Bloom syndrome
214400	Charcot-Marie-Tooth neuropathy-4A
216550	Cohen syndrome
216900	Achromatopsia
217000	C2 deficiency
217000	C3b inactivator deficiency
<u> </u>	Oso macrivator deficiency

217050	C6 deficiency
217050	Combined C6/C7 deficiency
217070	C7 deficiency
217095	Conotruncal cardiac anomalies
217300	Cornea plana congenita, recessive
217800	Macular corneal dystrophy
218000	Andermann syndrome
218030	Apparent mineralocorticoid excess, hypertension due to
221770	Polycystic lipomembranous osteodysplasia with sclerosing
	leukencephalopathy
221820	Gliosis, familial progressive subcortical
222100	Diabetes mellitus, insulin-dependent-1
222600	Atelosteogenesis II, 256050
222600	Achondrogenesis Ib, 600972
222600	Diastrophic dysplasia
222700	Lysinuric protein intolerance
222800	Hemolytic anemia due to bisphosphoglycerate mutase deficiency
222900	Sucrose intolerance
223900	Dysautonomia, familial
224100	Congenital dyserythropoietic anemia II
225500	Ellis-van Creveld syndrome
227220	[Eye color, brown]
227646	Fanconi anemia, type D
229300	Friedreich ataxia
229300	Friedreich ataxia with retained reflexes
229600	Fructose intolerance
229800	[Fructosuria]
230000	Fucosidosis
230350	Galactose epimerase deficiency
230400	Galactosemia
230800	Gaucher disease
230800	Gaucher disease with cardiovascular calcification
231550	Achalasia-addisonianism-alacrimia syndrome
231680	Glutaricaciduria, type IIA
232050	Propionicacidemia, type II or pccB type
232200	Glycogen storage disease I
232400	Glycogen storage disease IIIa
232400	Glycogen storage disease IIIb
233100	[Renal glucosuria]
233710	Chronic granulomatous disease due to deficiency of NCF-2
234000	Factor XII deficiency
235200	Hemochromatosis
235800	[Histidinemia]
236100	Holoprosencephaly-1
236200	Homocystinuria, B6-responsive and nonresponsive types
236730	Urofacial syndrome

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264300	Pseudohermaphroditism, male, with gynecomastia
264700	Pseudo-vitamin D dependency rickets 1
266200	Anemia, hemolytic, due to PK deficiency
266600	Inflammatory bowel disease-1
267750	Knobloch syndrome
268800	Sandhoff disease, infantile, juvenile, and adult forms
268800	Spinal muscular atrophy, HEXB-related
269920	Salla disease
270100	Situs inversus viscerum
271245	Spinocerebellar ataxia-8, infantile, with sensory neuropathy
271900	Canavan disease
272750	GM2-gangliosidosis, AB variant
272800	Tay-Sachs disease
272800	[Hex A pseudodeficiency]
272800	GM2-gangliosidosis, juvenile, adult
273300	Male germ cell tumor
274600	Pendred syndrome
274600	Deafness, autosomal recessive 4
276600	Tyrosinemia, type II
276700	Tyrosinemia, type I
276900	Usher syndrome, type 1A
276902	Usher syndrome, type 3
278000	Wolman disease
278000	Cholesteryl ester storage disease
278300	Xanthinuria, type I
278760	Xeroderma pigmentosum, group F
300031	Mental retardation, X-linked, FRAXF type
300044	Wernicke-Korsakoff syndrome, susceptibility to
300048	Intestinal pseudoobstruction, neuronal, X-linked
300049	Nodular heterotopia, bilateral periventricular
300049	BPNH/MR syndrome
300055	Mental retardation with psychosis, pyramidal signs, and
	macroorchidism
300100	Adrenoleukodystrophy
300100	Adrenomyeloneuropathy
300104	Mental retardation, X-linked nonspecific, 309541
300126	Dyskeratosis congenita-1, 305000
301201	Amelogenesis imperfecta-3, hypoplastic type
301220	Partington syndrome II
301590	Anophthalmos-1
302060	Noncompaction of left ventricular myocardium, isolated
302060	Barth syndrome
302060	Cardiomyopathy, X-linked dilated, 300069
302060	Endocardial fibroelastosis-2
302350	Nance-Horan syndrome
302960	Chondrodysplasia punctata, X-linked dominant

303700 Colorblindness, blue monochromatic		
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415000 Sertoli-cell-only syndrome		
600020 Prostate cancer, 176807		
	600020	Prostate cancer, 176807

Colomatal
Colorectal cancer
Myelodysplasia syndrome-l
Retinitis pigmentosa-13
Leukocyte adhesion deficiency, 116920
Split hand/foot malformation, type 3
Deafness, autosomal dominant 2
Retinitis pigmentosa-12, autosomal recessive
Muscular dystrophy, Duchenne-like, type 2
Adhalinopathy, primary
Retinitis pigmentosa-11
Long QT syndrome-3
SCID, autosomal recessive, T-negative/B-positive type
Spinal muscular atrophy, congenital nonprogressive, of lower
limbs
Leber congenital amaurosis, type I, 204000
Carnitine acetyltransferase deficiency
Ichthyosis bullosa of Siemens, 146800
Dyslexia, specific, 2
Cleidocranial dysplasia, 119600
Spinocerebellar ataxia-4
Palmoplantar keratoderma, Bothnia type
Temperature-sensitive apoptosis
Ehlers-Danlos-like syndrome
Cerebral arteriopathy with subcortical infarcts and
leukoencephalopathy, 125310
Non-insulin-dependent diabetes mellitus, 125853
MODY, type 1, 125850
Atrioventricular canal defect-1
Pseudoachondroplasia, 177170
Epiphyseal dysplasia, multiple 1, 132400
Insulin-dependent diabetes mellitus-5
Tibial muscular dystrophy
Spinal muscular atrophy-1, 253300
Spinal muscular atrophy-2, 253550
Spinal muscular atrophy-3, 253400
Bardet-Biedl syndrome 4
[Ii blood group, 110800]
Epilepsy, partial
Myopathy, congenital
Chondrosarcoma, extraskeletal myxoid
Atrial septal defect with atrioventricular conduction defects, 108900
Craniosynostosis, Adelaide type
Myopathy due to CPT II deficiency, 255110
Nivopatity due to Cr 1 in deficiency, 2001 to
CPT deficiency, hepatic, type II, 600649

600698	Salivary adenoma
600698	Uterine leiomyoma
600698	Lipoma Lipoma
600698	Lipomatosis, mutiple, 151900
600701	Lipoma
600722	Ceroid lipofuscinosis, neuronal, variant juvenile type, with
	granular osmiophilic deposits
600722	Ceroid lipofuscinosis, neuronal-1, infantile, 256730
600757	Orofacial cleft-3
600759	Alzheimer disease-4
600760	Pseudohypoaldosteronism, type I, 264350
600760	Liddle syndrome, 177200
600761	Pseudohypoaldosteronism, type I, 264350
600761	Liddle syndrome, 177200
600807	Bronchial asthma
600808	Enuresis, nocturnal, 2
600837	Hirschsprung disease, 142623
600839	Bartter syndrome, 241200
600850	Schizophrenia disorder-4
600856	Beckwith-Wiedemann syndrome, 130650
600881	Cataract, congenital, zonular, with sutural opacities
600882	Charcot-Marie-Tooth neuropathy-2B
600883	Diabetes mellitus, insulin-dependent, 8
600884	Cardiomyopathy, familial dilated 1B
600887	Endometrial carcinoma
600897	Cataract, zonular pulverulent-1, 116200
600899	Severe combined immunodeficiency, type I, 202500
600918	Cystinuria, type III
600919	Long QT syndrome-4 with sinus bradycardia
600923	Porphyria variegata, 176200
600946	Short stature, autosomal dominant, with normal serum growth
	hormone binding protein
600946	Short stature, idiopathic
600946	Laron dwarfism, 262500
600956	Persistent Mullerian duct syndrome, type II, 261550
600957	Persistent Mullerian duct syndrome, type I, 261550
600968	Gitelman syndrome, 263800
600971	Deafness, autosomal recessive 6
600974	Deafness, autosomal recessive 7
600977	Cone dystrophy, progressive
600994	Deafness, autosomal dominant 5
600995	Nephrotic syndrome, idiopathic, steroid-resistant
600998	Bleeding diathesis due to GNAQ deficiency
601002	5-oxoprolinuria, 266130
601002	Hemolytic anemia due to glutathione synthetase deficiency,
	231900
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601071	Deafness, autosomal recessive 9
601071	Deafness, autosomal recessive 8
601105	Pycnodysostosis, 265800
601107	Dubin-Johnson syndrome, 237500
601130	Tolbutamide poor metabolizer
601145	Epilepsy, progressive myoclonic 1, 254800
601146	Brachydactyly, type C, 113100
601146	Acromesomelic dysplasia, Hunter-Thompson type, 201250
601146	Chondrodysplasia, Grebe type, 200700
601154	Cardiomyopathy, dilated, 1E
601199	Neonatal hyperparathyroidism, 239200
601199	Hypocalcemia, autosomal dominant, 601198
601199	Hypocalciuric hypercalcemia, type I, 145980
601202	Cataract, anterior polar-2
601208	Insulin-dependent diabetes mellitus-11
601226	Progressive external ophthalmoplegia, type 2
601238	Cerebellar ataxia, Cayman type
601267	HIV infection, susceptibility/resistence to
601284	Hereditary hemorrhagic telangiectasia-2, 600376
601313	Polycystic kidney disease, adult type I, 173900
601316	Deafness, autosomal dominant 10
601363	Wilms tumor, type 4
601373	HIV infection, susceptibility/resistance to
601402	Leukemia, myeloid, acute
601406	B-cell non-Hodgkin lymphoma, high-grade
601410	Diabetes mellitus, transient neonatal
601411	Muscular dystrophy, limb-girdle, type 2F, 601287
601412	Deafness, autosomal dominant 7
601414	Retinitis pigmentosa-18
601471·	Moebius syndrome-2
601493	Cardiomyopathy, dilated 1C
601494	Cardiomyopathy, familial, dilated-2
601518	Prostate cancer, hereditary, 1, 176807
601542	Rieger syndrome, type 1, 180500
601545	Lissencephaly-1
601596	Charcot-Marie-Tooth neuropathy, demyelinating
601604	Mycobacterial and salmonella infections, susceptibility to
601607	Rhabdoid tumors
601620	Holt-Oram syndrome, 142900
601621	Ulnar-mammary syndrome, 181450
601649	Blepharophimosis, epicanthus inversus, and ptosis, type 2
601652	Glaucoma 1A, primary open angle, juvenile-onset, 137750
601669	Hirschsprung disease, one form
601680	Distal arthrogryposis, type 2B
601682	Glaucoma 1C, primary open angle
	Platelet-activating factor acetylhydrolase deficiency
601690	Transfer activating factor acceptify diotase deficiency

601691	Retinitis pigmentosa-19, 601718
601691	Stargardt disease-1, 248200
601691	Cone-rod dystrophy 3
601691	Fundus flavimaculatus with macular dystrophy, 248200
601718	Retinitis pigmentosa-19
601757	Rhizomelic chondrodysplasia punctata, type 1, 215100
601769	Osteoporosis, involutional
601769	Rickets, vitamin D-resistant, 277440
601771	Glaucoma 3A, primary infantile, 231300
601777	Cone dystrophy, progressive
601780	Ceroid-lipofuscinosis, neuronal-6, variant late infantile
601800	[Hair color, brown]
601841	Protein C inhibitor deficiency
601843	Hypothyroidism, congenital, 274400
601844	Pseudohypoaldosteronism type II
601846	Muscular dystrophy with rimmed vacuoles
601863	Bare lymphocyte syndrome, complementation group C
601868	Deafness, autosomal dominant 13
601885	Cataract, zonular pulverulent-2
601916	Pancreatic cancer
601928	Monilethrix, 158000
601954	Muscular dystrophy, limb-girdle, type 2G
601969	Glioblastoma multiforme, 137800
601969	Medulloblastoma, 155255
601975	Ectodermal dysplasia/skin fragility syndrome
602014	Hypomagnesemia with secondary hypocalcemia
602066	Convulsions, infantile and paroxysmal choreoathetosis
602067	Cardiomyopathy, dilated, 1F
602081	Speech-language disorder-1
602082	Corneal dystrophy, Thiel-Behnke type
602084	Endometrial carcinoma
602088	Nephronophthisis, infantile
602089	Hemangioma, capillary, hereditary
602092	Deafness, autosomal recessive 18
602094	Lipodystrophy, familial partial
602116	Glioma
602134	Tremor, familial essential, 2
602136	Refsum disease, infantile, 266510
602136	Zellweger syndrome-1, 214100
602136	Adrenoleukodystrophy, neonatal, 202370
602153	Monilethrix, 158000
602216	Peutz-Jeghers syndrome, 175200
602221	Stem-cell leukemia/lymphoma syndrome
602225	Cone-rod retinal dystrophy-2, 120970
602225	Leber congenital amaurosis, type III
602279	Oculopharyngeal muscular dystorphy, 164300
002213	Obusopharyngear museurar dysiorphy, 104300

602279	Oculopharyngeal muscular dystrophy, autosomal recessive, 257950
602280	Retinitis pigmentosa-14, 600132
602363	Ellis-van Creveld-like syndrome
602403	Alzheimer disease, susceptibility to
602421	Sweat chloride elevation without CF
602421	Congenital bilateral absence of vas deferens, 277180
602421	Cystic fibrosis, 219700
602447	Coronary artery disease, susceptibility to
602475	Ossification of posterior longitudinal ligament of spine
602476	Febrile convulsions, familial, 1
602477	Febrile convulsions, familial, 2
602491	Hyperlipidemia, familial combined, 1
602544	Parkinson disease, juvenile, type 2, 600116
602568	Homocystinuria-megaloblastic anemia, cbl E type, 236270
602574	Deafness, autosomal dominant 12, 601842
602574	Deafness, autosomal dominant 8, 601543
602575	Nail-patella syndrome with open-angle glaucoma, 137750
602575	Nail-patella syndrome, 161200
602629	Dystonia-6, torsion
602631	Rhabdomyosarcoma, 268210
602631	Breast Cancer
602669	Anterior segment mesenchymal dysgenesis and cataract, 107250
602669	Cataract, congenital
602685	Mental retardation, severe, with spasticity and tapetoretinal
	degeneration
602716	Nephrosis-1, congenital, Finnish type, 256300
602771	Muscular dystrophy, congenital, with early spine rigidity
602772	Retinitis pitmentosa-24

Polynucleotide and Polypeptide Variants

The present invention is also directed to variants of the immune/hematopoietic associated polynucleotide sequence disclosed in SEQ ID NO:X or the complementary strand thereto, nucleotide sequences encoding the polypeptide of SEQ ID NO:Y, the nucleotide sequence of SEQ ID NO:X encoding the polypeptide sequence as defined in column 6 of Table 1A, nucleotide sequences encoding the polypeptide as defined in column 6 of Table 1A, the nucleotide sequence as defined in columns 8 and 9 of Table 2, nucleotide sequences encoding the polypeptide encoded by the nucleotide sequence as defined in columns 8 and 9 of Table 2, the nucleotide sequence as defined in column 6 of Table 1B, nucleotide sequences encoding the polypeptide encoded by the nucleotide sequence as defined in column 6 of Table 1B, the cDNA sequence contained in Clone ID NO:Z, and/or nucleotide sequences encoding a polypeptide encoded by the cDNA sequence contained in Clone ID NO:Z.

- [092] The present invention also encompasses variants of the polypeptide sequence disclosed in SEQ ID NO:Y, a polypeptide sequence as defined in column 6 of Table 1A, a polypeptide sequence encoded by the polynucleotide sequence in SEQ ID NO:X, a polypeptide sequence encoded by the nucleotide sequence as defined in columns 8 and 9 of Table 2, a polypeptide sequence encoded by the nucleotide sequence as defined in column 6 of Table 1B, a polypeptide sequence encoded by the complement of the polynucleotide sequence in SEQ ID NO:X, and/or a polypeptide sequence encoded by the cDNA sequence contained in Clone ID NO:Z.
- [093] "Variant" refers to a polynucleotide or polypeptide differing from the polynucleotide or polypeptide of the present invention, but retaining essential properties thereof. Generally, variants are overall closely similar, and, in many regions, identical to the polynucleotide or polypeptide of the present invention.
- [094] Thus, one aspect of the invention provides an isolated nucleic acid molecule comprising, or alternatively consisting of, a polynucleotide having a nucleotide sequence selected from the group consisting of: (a) a nucleotide sequence described in SEQ ID NO:X or contained in the cDNA sequence of Clone ID NO:Z; (b) a nucleotide sequence in SEQ ID NO:X or the cDNA in Clone ID NO:Z which encodes a mature immune/hematopoietic associated polypeptide; (c) a nucleotide sequence in SEQ ID NO:X or the cDNA sequence of Clone ID NO:Z, which encodes a biologically active

fragment of an immune/hematopoietic associated polypeptide; (d) a nucleotide sequence in SEQ ID NO:X or the cDNA sequence of Clone ID NO:Z, which encodes an antigenic fragment of an immune/hematopoietic associated polypeptide; (e) a nucleotide sequence encoding an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; (f) a nucleotide sequence encoding a mature immune/hematopoietic associated polypeptide of the amino acid sequence of SEQ ID NO:Y or the amino acid sequence encoded by the cDNA in Clone ID NO:Z; (g) a nucleotide sequence encoding a biologically active fragment of an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; (h) a nucleotide sequence encoding an antigenic fragment of an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; and (i) a nucleotide sequence complementary to any of the nucleotide sequences in (a), (b), (c), (d), (e), (f), (g), or (h), above.

The present invention is also directed to nucleic acid molecules which [095] comprise, or alternatively consist of, a nucleotide sequence which is at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99% or 100%, identical to, for example, any of the nucleotide sequences in (a), (b), (c), (d), (e), (f), (g), (h), or (i) above, the nucleotide coding sequence in SEQ ID NO:X or the complementary strand thereto, the nucleotide coding sequence of the cDNA contained in Clone ID NO:Z or the complementary strand thereto, a nucleotide sequence encoding the polypeptide of SEQ ID NO:Y, a nucleotide sequence encoding a polypeptide sequence encoded by the nucleotide sequence in SEQ ID NO:X, a polypeptide sequence encoded by the complement of the polynucleotide sequence in SEQ ID NO:X, a nucleotide sequence encoding the polypeptide encoded by the cDNA contained in Clone ID NO:Z, the nucleotide coding sequence in SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or the complementary strand thereto, a nucleotide sequence encoding the polypeptide encoded by the nucleotide sequence in SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or the complementary strand thereto, the nucleotide coding sequence in SEQ ID NO:B as defined in column 6 of Table IB or the complementary strand thereto, a nucleotide sequence encoding the polypeptide encoded by the nucleotide sequence in SEQ

ID NO:B as defined in column 6 of Table 1B or the complementary strand thereto, the nucleotide sequence in SEQ ID NO:X encoding the polypeptide sequence as defined in column 6 of Table 1A or the complementary strand thereto, nucleotide sequences encoding a polypeptide as defined in column 6 of Table 1A or the complementary strand thereto, and/or polynucleotide fragments of any of these nucleic acid molecules (e.g., those fragments described herein). Polynucleotides which hybridize to the complement of these nucleic acid molecules under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention, as are polypeptides encoded by these polynucleotides and nucleic acids.

[096] In a preferred embodiment, the invention encompasses nucleic acid molecules which comprise, or alternatively, consist of a polynucleotide which hybridizes under stringent hybridization conditions, or alternatively, under lower stringency conditions, to a polynucleotide in (a), (b), (c), (d), (e), (f), (g), (h), or (i) above, as are polypeptides encoded by these polynucleotides. In another preferred embodiment, polynucleotides which hybridize to the complement of these nucleic acid molecules under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention, as are polypeptides encoded by these polynucleotides.

In another embodiment, the invention provides a purified protein comprising, or alternatively consisting of, a polypeptide having an amino acid sequence selected from the group consisting of: (a) the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; (b) the amino acid sequence of a mature immune/hematopoietic associated polypeptide having the amino acid sequence of SEQ ID NO:Y or the amino acid sequence encoded by the cDNA in Clone ID NO:Z; (c) the amino acid sequence of a biologically active fragment of an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z; and (d) the amino acid sequence of an antigenic fragment of an immune/hematopoietic associated polypeptide having the complete amino acid sequence of SEQ ID NO:Y or the complete amino acid sequence encoded by the cDNA in Clone ID NO:Z.

[098] The present invention is also directed to proteins which comprise, or alternatively consist of, an amino acid sequence which is at least 80%, 85%, 90%, 95%,

96%, 97%, 98%, 99% or 100%, identical to, for example, any of the amino acid sequences in (a), (b), (c), or (d), above, the amino acid sequence shown in SEQ ID NO:Y, the amino acid sequence encoded by the cDNA contained in Clone ID NO:Z, the amino acid sequence of the polypeptide encoded by the nucleotide sequence in SEQ ID NO:X as defined in columns 8 and 9 of Table 2, the amino acid sequence of the polypeptide encoded by the nucleotide sequence in SEQ ID NO:B as defined in column 6 of Table 1B, the amino acid sequence as defined in column 6 of Table 1A, an amino acid sequence encoded by the nucleotide sequence in SEQ ID NO:X, and an amino acid sequence encoded by the complement of the polynucleotide sequence in SEQ ID NO:X. Fragments of these polypeptides are also provided (e.g., those fragments described herein). Further proteins encoded by polynucleotides which hybridize to the complement of the nucleic acid molecules encoding these amino acid sequences under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention, as are the polynucleotides encoding these proteins.

[099] By a nucleic acid having a nucleotide sequence at least, for example, 95% "identical" to a reference nucleotide sequence of the present invention, it is intended that the nucleotide sequence of the nucleic acid is identical to the reference sequence except that the nucleotide sequence may include up to five point mutations per each 100 nucleotides of the reference nucleotide sequence encoding the polypeptide. In other words, to obtain a nucleic acid having a nucleotide sequence at least 95% identical to a reference nucleotide sequence, up to 5% of the nucleotides in the reference sequence may be deleted or substituted with another nucleotide, or a number of nucleotides up to 5% of the total nucleotides in the reference sequence may be inserted into the reference sequence. The query sequence may be an entire sequence referred to in Table 1A or 2 as the ORF (open reading frame), or any fragment specified, as described herein.

[0100] As a practical matter, whether any particular nucleic acid molecule or polypeptide is at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to a nucleotide sequence of the present invention can be determined conventionally using known computer programs. A preferred method for determining the best overall match between a query sequence (a sequence of the present invention) and a subject sequence, also referred to as a global sequence alignment, can be determined using the FASTDB computer program based on the algorithm of Brutlag et al. (Comp. App. Biosci. 6:237-245

(1990)). In a sequence alignment the query and subject sequences are both DNA sequences. An RNA sequence can be compared by converting U's to T's. The result of said global sequence alignment is expressed as percent identity. Preferred parameters used in a FASTDB alignment of DNA sequences to calculate percent identity are: Matrix=Unitary, k-tuple=4, Mismatch Penalty=1, Joining Penalty=30, Randomization Group Length=0, Cutoff Score=1, Gap Penalty=5, Gap Size Penalty 0.05, Window Size=500 or the length of the subject nucleotide sequence, whichever is shorter.

[0101] If the subject sequence is shorter than the query sequence because of 5' or 3' deletions, not because of internal deletions, a manual correction must be made to the results. This is because the FASTDB program does not account for 5' and 3' truncations of the subject sequence when calculating percent identity. For subject sequences truncated at the 5' or 3' ends, relative to the query sequence, the percent identity is corrected by calculating the number of bases of the query sequence that are 5' and 3' of the subject sequence, which are not matched/aligned, as a percent of the total bases of the query sequence. Whether a nucleotide is matched/aligned is determined by results of the FASTDB sequence alignment. This percentage is then subtracted from the percent identity, calculated by the above FASTDB program using the specified parameters, to arrive at a final percent identity score. This corrected score is what is used for the purposes of the present invention. Only bases outside the 5' and 3' bases of the subject sequence, as displayed by the FASTDB alignment, which are not matched/aligned with the query sequence, are calculated for the purposes of manually adjusting the percent identity score.

sequence to determine percent identity. The deletions occur at the 5' end of the subject sequence and therefore, the FASTDB alignment does not show a matched/alignment of the first 10 bases at 5' end. The 10 unpaired bases represent 10% of the sequence (number of bases at the 5' and 3' ends not matched/total number of bases in the query sequence) so 10% is subtracted from the percent identity score calculated by the FASTDB program. If the remaining 90 bases were perfectly matched the final percent identity would be 90%. In another example, a 90 base subject sequence is compared with a 100 base query sequence. This time the deletions are internal deletions so that there are no bases on the 5' or 3' of the subject sequence which are not matched/aligned with the query. In this case

the percent identity calculated by FASTDB is not manually corrected. Once again, only bases 5' and 3' of the subject sequence which are not matched/aligned with the query sequence are manually corrected for. No other manual corrections are to be made for the purposes of the present invention.

"identical" to a query amino acid sequence of the present invention, it is intended that the amino acid sequence of the subject polypeptide is identical to the query sequence except that the subject polypeptide sequence may include up to five amino acid alterations per each 100 amino acids of the query amino acid sequence. In other words, to obtain a polypeptide having an amino acid sequence at least 95% identical to a query amino acid sequence, up to 5% of the amino acid residues in the subject sequence may be inserted, deleted, (indels) or substituted with another amino acid. These alterations of the reference sequence may occur at the amino or carboxy terminal positions of the reference amino acid sequence or anywhere between those terminal positions, interspersed either individually among residues in the reference sequence or in one or more contiguous groups within the reference sequence.

[0104] As a practical matter, whether any particular polypeptide is at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to, for instance, the amino acid sequence of a polypeptide referred to in Table 1A (e.g., an amino acid sequence identified in columns 5 or 6) or Table 2 (e.g., the amino acid sequence of the polypeptide encoded by the polynucleotide sequence defined in columns 8 and 9 of Table 2) or a fragment thereof, the amino acid sequence of the polypeptide encoded by the polynucleotide sequence in SEQ ID NO:B as defined in column 6 of Table 1B or a fragment thereof, the amino acid sequence of the polypeptide encoded by the nucleotide sequence in SEQ ID NO:X or a fragment thereof, or an amino acid sequence of the polypeptide encoded by cDNA contained in Clone ID NO:Z, or a fragment thereof, can be determined conventionally using known computer programs. A preferred method for determining the best overall match between a query sequence (a sequence of the present invention) and a subject sequence, also referred to as a global sequence alignment, can be determined using the FASTDB computer program based on the algorithm of Brutlag et al. (Comp. App. Biosci.6:237-245 (1990)). In a sequence alignment the query and subject sequences are either both nucleotide sequences or both amino acid sequences. The result of said global

sequence alignment is expressed as percent identity. Preferred parameters used in a FASTDB amino acid alignment are: Matrix=PAM 0, k-tuple=2, Mismatch Penalty=1, Joining Penalty=20, Randomization Group Length=0, Cutoff Score=1, Window Size=sequence length, Gap Penalty=5, Gap Size Penalty=0.05, Window Size=500 or the length of the subject amino acid sequence, whichever is shorter.

[0105] If the subject sequence is shorter than the query sequence due to N- or Cterminal deletions, not because of internal deletions, a manual correction must be made to the results. This is because the FASTDB program does not account for N- and C-terminal truncations of the subject sequence when calculating global percent identity. For subject sequences truncated at the N- and C-termini, relative to the query sequence, the percent identity is corrected by calculating the number of residues of the query sequence that are N- and C-terminal of the subject sequence, which are not matched/aligned with a corresponding subject residue, as a percent of the total bases of the query sequence. Whether a residue is matched/aligned is determined by results of the FASTDB sequence alignment. This percentage is then subtracted from the percent identity, calculated by the above FASTDB program using the specified parameters, to arrive at a final percent identity score. This final percent identity score is what is used for the purposes of the present invention. Only residues to the N- and C-termini of the subject sequence, which are not matched/aligned with the query sequence, are considered for the purposes of manually adjusting the percent identity score. That is, only query residue positions outside the farthest N- and C- terminal residues of the subject sequence.

residue query sequence to determine percent identity. The deletion occurs at the N-terminus of the subject sequence and therefore, the FASTDB alignment does not show a matching/alignment of the first 10 residues at the N-terminus. The 10 unpaired residues represent 10% of the sequence (number of residues at the N- and C- termini not matched/total number of residues in the query sequence) so 10% is subtracted from the percent identity score calculated by the FASTDB program. If the remaining 90 residues were perfectly matched the final percent identity would be 90%. In another example, a 90 residue subject sequence is compared with a 100 residue query sequence. This time the deletions are internal deletions so there are no residues at the N- or C-termini of the subject sequence which are not matched/aligned with the query. In this case the percent

identity calculated by FASTDB is not manually corrected. Once again, only residue positions outside the N- and C-terminal ends of the subject sequence, as displayed in the FASTDB alignment, which are not matched/aligned with the query sequence are manually corrected for. No other manual corrections are to be made for the purposes of the present invention.

[0107] The polynucleotide variants of the invention may contain alterations in the coding regions, non-coding regions, or both. Especially preferred are polynucleotide variants containing alterations which produce silent substitutions, additions, or deletions, but do not alter the properties or activities of the encoded polypeptide. Nucleotide variants produced by silent substitutions due to the degeneracy of the genetic code are preferred. Moreover, polypeptide variants in which less than 50, less than 40, less than 30, less than 20, less than 10, or 5-50, 5-25, 5-10, 1-5, or 1-2 amino acids are substituted, deleted, or added in any combination are also preferred. Polynucleotide variants can be produced for a variety of reasons, e.g., to optimize codon expression for a particular host (change codons in the human mRNA to those preferred by a bacterial host such as *E. coli*).

[0108] Naturally occurring variants are called "allelic variants," and refer to one of several alternate forms of a gene occupying a given locus on a chromosome of an organism. (Genes II, Lewin, B., ed., John Wiley & Sons, New York (1985).) These allelic variants can vary at either the polynucleotide and/or polypeptide level and are included in the present invention. Alternatively, non-naturally occurring variants may be produced by mutagenesis techniques or by direct synthesis.

[0109] Using known methods of protein engineering and recombinant DNA technology, variants may be generated to improve or alter the characteristics of the polypeptides of the present invention. For instance, one or more amino acids can be deleted from the N-terminus or C-terminus of the polypeptides of the present invention without substantial loss of biological function. As an example, the authors of Ron et al., J. Biol. Chem. 268: 2984-2988 (1993), reported variant KGF proteins having heparin binding activity even after deleting 3, 8, or 27 amino-terminal amino acid residues. Similarly, Interferon gamma exhibited up to ten times higher activity after deleting 8-10 amino acid residues from the carboxy terminus of this protein. (Dobeli et al., J. Biotechnology 7:199-216 (1988).)

[0110] Moreover, ample evidence demonstrates that variants often retain a biological activity similar to that of the naturally occurring protein. For example, Gayle and coworkers (J. Biol. Chem. 268:22105-22111 (1993)) conducted extensive mutational analysis of human cytokine IL-1a. They used random mutagenesis to generate over 3,500 individual IL-1a mutants that averaged 2.5 amino acid changes per variant over the entire length of the molecule. Multiple mutations were examined at every possible amino acid position. The investigators found that "[m]ost of the molecule could be altered with little effect on either [binding or biological activity]." In fact, only 23 unique amino acid sequences, out of more than 3,500 nucleotide sequences examined, produced a protein that significantly differed in activity from wild-type.

- [0111] Furthermore, even if deleting one or more amino acids from the N-terminus or C-terminus of a polypeptide results in modification or loss of one or more biological functions, other biological activities may still be retained. For example, the ability of a deletion variant to induce and/or to bind antibodies which recognize the secreted form will likely be retained when less than the majority of the residues of the secreted form are removed from the N-terminus or C-terminus. Whether a particular polypeptide lacking N-or C-terminal residues of a protein retains such immunogenic activities can readily be determined by routine methods described herein and otherwise known in the art.
- [0112] Thus, the invention further includes polypeptide variants which show a functional activity (e.g., biological activity) of the polypeptides of the invention. Such variants include deletions, insertions, inversions, repeats, and substitutions selected according to general rules known in the art so as have little effect on activity.
- [0113] The present application is directed to nucleic acid molecules at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99% or 100% identical to the nucleic acid sequences disclosed herein, (e.g., encoding a polypeptide having the amino acid sequence of an N and/or C terminal deletion), irrespective of whether they encode a polypeptide having functional activity. This is because even where a particular nucleic acid molecule does not encode a polypeptide having functional activity, one of skill in the art would still know how to use the nucleic acid molecule, for instance, as a hybridization probe or a polymerase chain reaction (PCR) primer. Uses of the nucleic acid molecules of the present invention that do not encode a polypeptide having functional activity include, inter alia, (1) isolating a gene or allelic or splice variants thereof in a cDNA library; (2) in situ

hybridization (e.g., "FISH") to metaphase chromosomal spreads to provide precise chromosomal location of the gene, as described in Verma et al., Human Chromosomes: A Manual of Basic Techniques, Pergamon Press, New York (1988); (3) Northern Blot analysis for detecting mRNA expression in specific tissues (e.g., normal immune/hematopoietic or diseased immune/hematopoietic tissues); and (4) in situ hybridization (e.g., histochemistry) for detecting mRNA expression in specific tissues (e.g., normal immune/hematopoietic or diseased immune/hematopoietic tissues).

[0114] Preferred, however, are nucleic acid molecules having sequences at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99% or 100% identical to the nucleic acid sequences disclosed herein, which do, in fact, encode a polypeptide having functional activity. By a polypeptide having "functional activity" is meant, a polypeptide capable of displaying one or more known functional activities associated with a full-length (complete) protein of the invention. Such functional activities include, but are not limited to, biological activity, antigenicity [ability to bind (or compete with a polypeptide of the invention for binding) to an anti-polypeptide of the invention antibody], immunogenicity (ability to generate antibody which binds to a specific polypeptide of the invention), ability to form multimers with polypeptides of the invention, and ability to bind to a receptor or ligand for a polypeptide of the invention.

[0115] The functional activity of the polypeptides, and fragments, variants and derivatives of the invention, can be assayed by various methods.

[0116] For example, in one embodiment where one is assaying for the ability to bind or compete with full-length polypeptide of the present invention for binding to an antipolypeptide of the invention antibody, various immunoassays known in the art can be used, including but not limited to, competitive and non-competitive assay systems using techniques such as radioimmunoassays, ELISA (enzyme linked immunosorbent assay), "sandwich" immunoassays, immunoradiometric assays, gel diffusion precipitation reactions, immunodiffusion assays, in situ immunoassays (using colloidal gold, enzyme or radioisotope labels, for example), western blots, precipitation reactions, agglutination assays (e.g., gel agglutination assays, hemagglutination assays), complement fixation assays, immunofluorescence assays, protein A assays, and immunoelectrophoresis assays, etc. In one embodiment, antibody binding is detected by detecting a label on the primary antibody. In another embodiment, the primary antibody is detected by detecting binding

of a secondary antibody or reagent to the primary antibody. In a further embodiment, the secondary antibody is labeled. Many means are known in the art for detecting binding in an immunoassay and are within the scope of the present invention.

[0117] In another embodiment, where a ligand is identified, or the ability of a polypeptide fragment, variant or derivative of the invention to multimerize is being evaluated, binding can be assayed, e.g., by means well-known in the art, such as, for example, reducing and non-reducing gel chromatography, protein affinity chromatography, and affinity blotting. See generally, Phizicky et al., Microbiol. Rev. 59:94-123 (1995). In another embodiment, the ability of physiological correlates of a polypeptide of the present invention to bind to a substrate(s) of the polypeptide of the invention can be routinely assayed using techniques known in the art.

[0118] In addition, assays described herein (see Examples) and otherwise known in the art may routinely be applied to measure the ability of polypeptides of the present invention and fragments, variants and derivatives thereof to elicit polypeptide related biological activity (either *in vitro* or *in vivo*). Other methods will be known to the skilled artisan and are within the scope of the invention.

Of course, due to the degeneracy of the genetic code, one of ordinary skill in [0119] the art will immediately recognize that a large number of the nucleic acid molecules having a sequence at least 80%, 85%, 90%, 95%, 96%, 97%, 98%, 99%, or 100% identical to, for example, the nucleic acid sequence of the cDNA contained in Clone ID NO:Z, a nucleic acid sequence referred to in Table 1A (e.g., SEQ ID NO:X), a nucleic acid sequence disclosed in Table 2 (e.g., the nucleic acid sequence delineated in columns 8 and 9) or fragments thereof, will encode polypeptides "having functional activity." In fact, since degenerate variants of any of these nucleotide sequences all encode the same polypeptide, in many instances, this will be clear to the skilled artisan even without performing the above described comparison assay. It will be further recognized in the art that, for such nucleic acid molecules that are not degenerate variants, a reasonable number will also encode a polypeptide having functional activity. This is because the skilled artisan is fully aware of amino acid substitutions that are either less likely or not likely to significantly effect protein function (e.g., replacing one aliphatic amino acid with a second aliphatic amino acid), as further described below.

[0120] For example, guidance concerning how to make phenotypically silent amino acid substitutions is provided in Bowie et al., "Deciphering the Message in Protein Sequences: Tolerance to Amino Acid Substitutions," Science 247:1306-1310 (1990), wherein the authors indicate that there are two main strategies for studying the tolerance of an amino acid sequence to change.

- [0121] The first strategy exploits the tolerance of amino acid substitutions by natural selection during the process of evolution. By comparing amino acid sequences in different species, conserved amino acids can be identified. These conserved amino acids are likely important for protein function. In contrast, the amino acid positions where substitutions have been tolerated by natural selection indicates that these positions are not critical for protein function. Thus, positions tolerating amino acid substitution could be modified while still maintaining biological activity of the protein.
- [0122] The second strategy uses genetic engineering to introduce amino acid changes at specific positions of a cloned gene to identify regions critical for protein function. For example, site directed mutagenesis or alanine-scanning mutagenesis (introduction of single alanine mutations at every residue in the molecule) can be used. See Cunningham et al., Science 244:1081-1085 (1989). The resulting mutant molecules can then be tested for biological activity.
- [0123] As the authors state, these two strategies have revealed that proteins are surprisingly tolerant of amino acid substitutions. The authors further indicate which amino acid changes are likely to be permissive at certain amino acid positions in the protein. For example, most buried (within the tertiary structure of the protein) amino acid residues require nonpolar side chains, whereas few features of surface side chains are generally conserved. Moreover, tolerated conservative amino acid substitutions involve replacement of the aliphatic or hydrophobic amino acids Ala, Val, Leu and Ile; replacement of the hydroxyl residues Ser and Thr; replacement of the acidic residues Asp and Glu; replacement of the amide residues Asn and Gln, replacement of the basic residues Lys, Arg, and His; replacement of the aromatic residues Phe, Tyr, and Trp, and replacement of the small-sized amino acids Ala, Ser, Thr, Met, and Gly. Besides conservative amino acid substitutions, variants of the present invention include (i) substitutions with one or more of the non-conserved amino acid residues, where the substituted amino acid residues may or may not be one encoded by the genetic code, or (ii)

substitutions with one or more of the amino acid residues having a substituent group, or (iii) fusion of the mature polypeptide with another compound, such as a compound to increase the stability and/or solubility of the polypeptide (for example, polyethylene glycol), or (iv) fusion of the polypeptide with additional amino acids, such as, for example, an IgG Fc fusion region peptide, serum albumin (preferably human serum albumin) or a fragment or variant thereof, or leader or secretory sequence, or a sequence facilitating purification. Such variant polypeptides are deemed to be within the scope of those skilled in the art from the teachings herein.

- [0124] For example, polypeptide variants containing amino acid substitutions of charged amino acids with other charged or neutral amino acids may produce proteins with improved characteristics, such as less aggregation. Aggregation of pharmaceutical formulations both reduces activity and increases clearance due to the aggregate's immunogenic activity. See Pinckard et al., Clin. Exp. Immunol. 2:331-340 (1967); Robbins et al., Diabetes 36: 838-845 (1987); Cleland et al., Crit. Rev. Therapeutic Drug Carrier Systems 10:307-377 (1993).
- the amino acid sequence of a polypeptide having an amino acid sequence which comprise the amino acid sequence of a polypeptide having an amino acid sequence which contains at least one amino acid substitution, but not more than 50 amino acid substitutions, even more preferably, not more than 40 amino acid substitutions, still more preferably, not more than 30 amino acid substitutions, and still even more preferably, not more than 20 amino acid substitutions from a polypeptide sequence disclosed herein. Of course it is highly preferable for a polypeptide to have an amino acid sequence which comprises the amino acid sequence of a polypeptide of SEQ ID NO:Y, an amino acid sequence encoded by SEQ ID NO:X, an amino acid sequence encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, an amino acid sequence encoded by the complement of SEQ ID NO:X, and/or the amino acid sequence encoded by cDNA contained in Clone ID NO:Z which contains, in order of ever-increasing preference, at least one, but not more than 10, 9, 8, 7, 6, 5, 4, 3, 2 or 1 amino acid substitutions.
- [0126] In specific embodiments, the polypeptides of the invention comprise, or alternatively, consist of, fragments or variants of a reference amino acid sequence selected from: (a) the amino acid sequence of SEQ ID NO:Y or fragments thereof (e.g., the mature form and/or other fragments described herein); (b) the amino acid sequence encoded by

SEQ ID NO:X or fragments thereof; (c) the amino acid sequence encoded by the complement of SEQ ID NO:X or fragments thereof; (d) the amino acid sequence encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or fragments thereof; and (e) the amino acid sequence encoded by cDNA contained in Clone ID NO:Z or fragments thereof; wherein the fragments or variants have 1-5, 5-10, 5-25, 5-50, 10-50 or 50-150, amino acid residue additions, substitutions, and/or deletions when compared to the reference amino acid sequence. In preferred embodiments, the amino acid substitutions are conservative. Polynucleotides encoding these polypeptides are also encompassed by the invention.

Polynucleotide and Polypeptide Fragments

[0127] The present invention is also directed to polynucleotide fragments of the polynucleotides (nucleic acids) of the invention. In the present invention, a "polynucleotide fragment" refers to a polynucleotide having a nucleic acid sequence which, for example: is a portion of the cDNA contained in Clone ID NO:Z or the complementary strand thereto; is a portion of the polynucleotide sequence encoding the polypeptide encoded by the cDNA contained in Clone ID NO:Z or the complementary strand thereto; is a portion of a polynucleotide sequence encoding the amino acid sequence encoded by the region of SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or the complementary strand thereto; is a portion of the polynucleotide sequence of SEQ ID NO:X as defined in columns 8 and 9 of Table 2 or the complementary strand thereto; is a portion of the polynucleotide sequence in SEQ ID NO:X or the complementary strand thereto; is a polynucleotide sequence encoding a portion of the polypeptide of SEQ ID NO:Y; is a polynucleotide sequence encoding a portion of a polypeptide encoded by SEQ ID NO:X; is a polynucleotide sequence encoding a portion of a polypeptide encoded by the complement of the polynucleotide sequence in SEQ ID NO:X; is a portion of a polynucleotide sequence encoding the amino acid sequence encoded by the region of SEQ ID NO:B as defined in column 6 of Table 1B or the complementary strand thereto; or is a portion of the polynucleotide sequence of SEQ ID NO:B as defined in column 6 of Table 1B or the complementary strand thereto.

[0128] The polynucleotide fragments of the invention are preferably at least about 15 nt, and more preferably at least about 20 nt, still more preferably at least about 30 nt, and

even more preferably, at least about 40 nt, at least about 50 nt, at least about 75 nt, or at least about 150 nt in length. A fragment "at least 20 nt in length," for example, is intended to include 20 or more contiguous bases from the cDNA sequence contained in Clone ID NO:Z, or the nucleotide sequence shown in SEQ ID NO:X or the complementary stand thereto. In this context "about" includes the particularly recited value or a value larger or smaller by several (5, 4, 3, 2, or 1) nucleotides, at either terminus or at both termini. These nucleotide fragments have uses that include, but are not limited to, as diagnostic probes and primers as discussed herein. Of course, larger fragments (e.g., at least 160, 170, 180, 190, 200, 250, 500, 600, 1000, or 2000 nucleotides in length) are also encompassed by the invention.

Moreover, representative examples of polynucleotide fragments of the [0129] invention, comprise, or alternatively consist of, a sequence from about nucleotide number 1-50, 51-100, 101-150, 151-200, 201-250, 251-300, 301-350, 351-400, 401-450, 451-500, 501-550, 551-600, 651-700, 701-750, 751-800, 800-850, 851-900, 901-950, 951-1000, 1001-1050, 1051-1100, 1101-1150, 1151-1200, 1201-1250, 1251-1300, 1301-1350, 1351-1400, 1401-1450, 1451-1500, 1501-1550, 1551-1600, 1601-1650, 1651-1700, 1701-1750, 1751-1800, 1801-1850, 1851-1900, 1901-1950, 1951-2000, 2001-2050, 2051-2100, 2101-2150, 2151-2200, 2201-2250, 2251-2300, 2301-2350, 2351-2400, 2401-2450, 2451-2500, 2501-2550, 2551-2600, 2601-2650, 2651-2700, 2701-2750, 2751-2800, 2801-2850, 2851-2900, 2901-2950, 2951-3000, 3001-3050, 3051-3100, 3101-3150, 3151-3200, 3201-3250, 3251-3300, 3301-3350, 3351-3400, 3401-3450, 3451-3500, 3501-3550, 3551-3600, 3601-3650, 3651-3700, 3701-3750, 3751-3800, 3801-3850, 3851-3900, 3901-3950, 3951-4000, 4001-4050, 4051-4100, 4101-4150, 4151-4200, 4201-4250, 4251-4300, 4301-4350, 4351-4400, 4401-4450, 4451-4500, 4501-4550, 4551-4600, 4601-4650, 4651-4700, 4701-4750, 4751-4800, 4801-4850, 4851-4900, 4901-4950, 4951-5000, 5001-5050, 5051-5100, 5101-5150, 5151-5200, 5201-5250, 5251-5300, 5301-5350, 5351-5400, 5401-5450, 5451-5500, 5501-5550, 5551-5600, 5601-5650, 5651-5700, 5701-5750, 5751-5800, 5801-5850, 5851-5900, 5901-5950, 5951-6000, 6001-6050, 6051-6100, 6101-6150, 6151-6200, 6201-6250, 6251-6300, 6301-6350, 6351-6400, 6401-6450, 6451-6500, 6501-6550, 6551-6600, 6601-6650, 6651-6700, 6701-6750, 6751-6800, 6801-6850, 6851-6900, 6901-6950, 6951-7000, 7001-7050, 7051-7100, 7101-7150, 7151-7200, 7201-7250, 7251-7300 or 7301 to the end of SEQ ID NO:X, or the complementary strand thereto. In this context "about" includes

the particularly recited range or a range larger or smaller by several (5, 4, 3, 2, or 1) nucleotides, at either terminus or at both termini. Preferably, these fragments encode a polypeptide which has a functional activity (e.g., biological activity). More preferably, these polynucleotides can be used as probes or primers as discussed herein. Polynucleotides which hybridize to one or more of these polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions are also encompassed by the invention, as are polypeptides encoded by these polynucleotides.

[0130] Further representative examples of polynucleotide fragments of the invention, comprise, or alternatively consist of, a sequence from about nucleotide number 1-50, 51-100, 101-150, 151-200, 201-250, 251-300, 301-350, 351-400, 401-450, 451-500, 501-550, 551-600, 651-700, 701-750, 751-800, 800-850, 851-900, 901-950, 951-1000, 1001-1050, 1051-1100, 1101-1150, 1151-1200, 1201-1250, 1251-1300, 1301-1350, 1351-1400, 1401-1450, 1451-1500, 1501-1550, 1551-1600, 1601-1650, 1651-1700, 1701-1750, 1751-1800, 1801-1850, 1851-1900, 1901-1950, 1951-2000, 2001-2050, 2051-2100, 2101-2150, 2151-2200, 2201-2250, 2251-2300, 2301-2350, 2351-2400, 2401-2450, 2451-2500, 2501-2550, 2551-2600, 2601-2650, 2651-2700, 2701-2750, 2751-2800, 2801-2850, 2851-2900, 2901-2950, 2951-3000, 3001-3050, 3051-3100, 3101-3150, 3151-3200, 3201-3250, 3251-3300, 3301-3350, 3351-3400, 3401-3450, 3451-3500, 3501-3550, 3551-3600, 3601-3650, 3651-3700, 3701-3750, 3751-3800, 3801-3850, 3851-3900, 3901-3950, 3951-4000, 4001-4050, 4051-4100, 4101-4150, 4151-4200, 4201-4250, 4251-4300, 4301-4350, 4351-4400, 4401-4450, 4451-4500, 4501-4550, 4551-4600, 4601-4650, 4651-4700, 4701-4750, 4751-4800, 4801-4850, 4851-4900, 4901-4950, 4951-5000, 5001-5050, 5051-5100, 5101-5150, 5151-5200, 5201-5250, 5251-5300, 5301-5350, 5351-5400, 5401-5450, 5451-5500, 5501-5550, 5551-5600, 5601-5650, 5651-5700, 5701-5750, 5751-5800, 5801-5850, 5851-5900, 5901-5950, 5951-6000, 6001-6050, 6051-6100, 6101-6150, 6151-6200, 6201-6250, 6251-6300, 6301-6350, 6351-6400, 6401-6450, 6451-6500, 6501-6550, 6551-6600, 6601-6650, 6651-6700, 6701-6750, 6751-6800, 6801-6850, 6851-6900, 6901-6950, 6951-7000, 7001-7050, 7051-7100, 7101-7150, 7151-7200, 7201-7250, 7251-7300 or 7301 to the end of the cDNA sequence contained in Clone ID NO:Z, or the complementary strand thereto. In this context "about" includes the particularly recited range or a range larger or smaller by several (5, 4, 3, 2, or 1) nucleotides, at either terminus or at both termini. Preferably, these fragments encode a polypeptide which has a functional activity (e.g., biological activity).

More preferably, these polynucleotides can be used as probes or primers as discussed herein. Polynucleotides which hybridize to one or more of these polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions are also encompassed by the invention, as are polypeptides encoded by these polynucleotides.

[0131] Moreover, representative examples of polynucleotide fragments of the invention comprise, or alternatively consist of, a nucleic acid sequence comprising one, two, three, four, five, six, seven, eight, nine, ten, or more of the above described polynucleotide fragments of the invention in combination with a polynucleotide sequence delineated in Table 1B column 6. Additional, representative examples of polynucleotide fragments of the invention comprise, or alternatively consist of, a nucleic acid sequence comprising one, two, three, four, five, six, seven, eight, nine, ten, or more of the above described polynucleotide fragments of the invention in combination with a polynucleotide sequence that is the complementary strand of a sequence delineated in column 6 of Table In further embodiments, the above-described polynucleotide fragments of the invention comprise, or alternatively consist of, sequences delineated in Table 1B, column 6, and have a nucleic acid sequence which is different from that of the BAC fragment having the sequence disclosed in SEQ ID NO:B (see Table 1B, column 5). In additional embodiments, the above-described polynucleotide fragments of the invention comprise, or alternatively consist of, sequences delineated in Table 1B, column 6, and have a nucleic acid sequence which is different from that published for the BAC clone identified as BAC ID NO:A (see Table 1B, column 4). In additional embodiments, the above-described polynucleotides of the invention comprise, or alternatively consist of, sequences delineated Table 1B, column 6, and have a nucleic acid sequence which is different from that contained in the BAC clone identified as BAC ID NO:A (see Table 1B, column 4). Polypeptides encoded by these polynucleotides, other polynucleotides that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides and polypeptides are also encompassed by the invention.

[0132] In additional specific embodiments, polynucleotides of the invention comprise, or alternatively consist of, one, two, three, four, five, six, seven, eight, nine, ten, or more fragments of the sequences delineated in column 6 of Table 1B, and the polynucleotide sequence of SEQ ID NO:X (e.g., as defined in Table 1B, column 2) or fragments or

variants thereof. Polypeptides encoded by these polynucleotides, other polynucleotides that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention.

- In additional specific embodiments, polynucleotides of the invention comprise, or alternatively consist of, one, two, three, four, five, six, seven, eight, nine, ten, or more fragments of the sequences delineated in column 6 of Table 1B which correspond to the same Clone ID NO:Z (see Table 1B, column 1), and the polynucleotide sequence of SEQ ID NO:X (e.g., as defined in Table 1A or 1B) or fragments or variants thereof. Polypeptides encoded by these polynucleotides, other polynucleotides that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention.
- In further specific embodiments, polynucleotides of the invention comprise, or alternatively consist of, one, two, three, four, five, six, seven, eight, nine, ten, or more fragments of the sequences delineated in the same row of column 6 of Table 1B, and the polynucleotide sequence of SEQ ID NO:X (e.g., as defined in Table 1A or 1B) or fragments or variants thereof. Polypeptides encoded by these polynucleotides, other polynucleotides that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention.
- [0135] In additional specific embodiments, polynucleotides of the invention comprise, or alternatively consist of a polynucleotide sequence in which the 3' 10 polynucleotides of one of the sequences delineated in column 6 of Table 1B and the 5' 10 polynucleotides of the sequence of SEQ ID NO:X are directly contiguous. Nucleic acids which hybridize to the complement of these 20 contiguous polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention. Polypeptides encoded by these polynucleotides and/or nucleic acids, other polynucleotides and/or nucleic acids that encode these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides, nucleic acids, and polypeptides are also encompassed by the invention.
- [0136] In additional specific embodiments, polynucleotides of the invention comprise, or alternatively consist of a polynucleotide sequence in which the 3' 10 polynucleotides of one of the sequences delineated in column 6 of Table 1B and the 5' 10 polynucleotides of

a fragment or variant of the sequence of SEQ ID NO:X (e.g., as described herein) are directly contiguous Nucleic acids which hybridize to the complement of these 20 contiguous polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention. Polypeptides encoded by these polynucleotides and/or nucleic acids, other polynucleotides and/or nucleic acids encoding these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides, nucleic acids, and polypeptides are also encompassed by the invention.

[0137] In further specific embodiments, polynucleotides of the invention comprise, or alternatively consist of a polynucleotide sequence in which the 3' 10 polynucleotides of a fragment or variant of the sequence of SEQ ID NO:X and the 5' 10 polynucleotides of the sequence of one of the sequences delineated in column 6 of Table 1B are directly contiguous. Nucleic acids which hybridize to the complement of these 20 contiguous polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention. Polypeptides encoded by these polynucleotides and/or nucleic acids, other polynucleotides and/or nucleic acids encoding these polypeptides, and antibodies that bind these polypeptides are also encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides, nucleic acids, and polypeptides are also encompassed by the invention.

[0138] In specific embodiments, polynucleotides of the invention comprise, or alternatively consist of a polynucleotide sequence in which the 3' 10 polynucleotides of one of the sequences delineated in column 6 of Table 1B and the 5' 10 polynucleotides of another sequence in column 6 are directly contiguous. In preferred embodiments, the 3' 10 polynucleotides of one of the sequences delineated in column 6 of Table 1B is directly contiguous with the 5' 10 polynucleotides of the next sequential exon delineated in Table 1B, column 6. Nucleic acids which hybridize to the complement of these 20 contiguous polynucleotides under stringent hybridization conditions or alternatively, under lower stringency conditions, are also encompassed by the invention. Polypeptides encoded by these polynucleotides and/or nucleic acids, other polynucleotides and/or nucleic acids encoding these polypeptides, and antibodies that bind these polypeptides are also

encompassed by the invention. Additionally, fragments and variants of the above-described polynucleotides, nucleic acids, and polypeptides are also encompassed by the invention.

In the present invention, a "polypeptide fragment" refers to an amino acid [0139] sequence which is a portion of that contained in SEQ ID NO:Y, a portion of an amino acid sequence encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, a portion of an amino acid sequence encoded by the polynucleotide sequence of SEQ ID NO:X, a portion of an amino acid sequence encoded by the complement of the polynucleotide sequence in SEQ ID NO:X, and/or a portion of an amino acid sequence encoded by the cDNA contained in Clone ID NO:Z. Protein (polypeptide) fragments may be "free-standing," or comprised within a larger polypeptide of which the fragment forms a part or region, most preferably as a single continuous region. Representative examples of polypeptide fragments of the invention, include, for example, fragments comprising, or alternatively consisting of, from about amino acid number 1-20, 21-40, 41-60, 61-80, 81-100, 102-120, 121-140, 141-160, 161-180, 181-200, 201-220, 221-240, 241-260, 261-280, 281-300, 301-320, 321-340, 341-360, 361-380, 381-400, 401-420, 421-440, 441-460, 461-480, 481-500, 501-520, 521-540, 541-560, 561-580, 581-600, 601-620, 621-640, 641-660, 661-680, 681-700, 701-720, 721-740, 741-760, 761-780, 781-800, 801-820, 821-840, 841-860, 861-880, 881-900, 901-920, 921-940, 941-960, 961-980, 981-1000, 1001-1020, 1021-1040, 1041-1060, 1061-1080, 1081-1100, 1101-1120, 1121-1140, 1141-1160, 1161-1180, 1181-1200, 1201-1220, 1221-1240, 1241-1260, 1261-1280, 1281-1300, 1301-1320, 1321-1340, 1341-1360, 1361-1380, 1381-1400, 1401-1420, 1421-1440, or 1441 to the end of the coding region. In a preferred embodiment, polypeptide fragments of the invention include, for example, fragments comprising, or alternatively consisting of, from about amino acid number 1-20, 21-40, 41-60, 61-80, 81-100, 102-120, 121-140, 141-160, 161-180, 181-200, 201-220, 221-240, 241-260, 261-280, 281-300, 301-320, 321-340, 341-360, 361-380, 381-400, 401-420, 421-440, 441-460, 461-480, 481-500, 501-520, 521-540, 541-560, 561-580, 581-600, 601-620, 621-640, 641-660, 661-680, 681-700, 701-720, 721-740, 741-760, 761-780, 781-800, 801-820, 821-840, 841-860, 861-880, 881-900, 901-920, 921-940, 941-960, 961-980, 981-1000, 1001-1020, 1021-1040, 1041-1060, 1061-1080, 1081-1100, 1101-1120, 1121-1140, 1141-1160, 1161-1180, 1181-1200, 1201-1220, 1221-1240, 1241-1260, 1261-1280, 1281-1300, 1301-1320, 1321-1340, 1341-1360, 1361-1380, 1381-

1400, 1401-1420, 1421-1440, or 1441 to the end of the coding region of SEQ ID NO:Y. Moreover, polypeptide fragments of the invention may be at least about 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 100, 110, 120, 130, 140, or 150 amino acids in length. In this context "about" includes the particularly recited ranges or values, or ranges or values larger or smaller by several (5, 4, 3, 2, or 1) amino acids, at either extreme or at both extremes. Polynucleotides encoding these polypeptide fragments are also encompassed by the invention.

- [0140] Even if deletion of one or more amino acids from the N-terminus of a protein results in modification of loss of one or more biological functions of the protein, other functional activities (e.g., biological activities, ability to multimerize, ability to bind a ligand) may still be retained. For example, the ability of shortened muteins to induce and/or bind to antibodies which recognize the complete or mature forms of the polypeptides generally will be retained when less than the majority of the residues of the complete or mature polypeptide are removed from the N-terminus. Whether a particular polypeptide lacking N-terminal residues of a complete polypeptide retains such immunologic activities can readily be determined by routine methods described herein and otherwise known in the art. It is not unlikely that a mutein with a large number of deleted N-terminal amino acid residues may retain some biological or immunogenic activities. In fact, peptides composed of as few as six amino acid residues may often evoke an immune response.
- [0141] Accordingly, polypeptide fragments include the secreted protein as well as the mature form. Further preferred polypeptide fragments include the secreted protein or the mature form having a continuous series of deleted residues from the amino or the carboxy terminus, or both. For example, any number of amino acids, ranging from 1-60, can be deleted from the amino terminus of either the secreted polypeptide or the mature form. Similarly, any number of amino acids, ranging from 1-30, can be deleted from the carboxy terminus of the secreted protein or mature form. Furthermore, any combination of the above amino and carboxy terminus deletions is preferred. Similarly, polynucleotides encoding these polypeptide fragments are also preferred.
- [0142] The present invention further provides polypeptides having one or more residues deleted from the amino terminus of the amino acid sequence of a polypeptide disclosed herein (e.g., a polypeptide of SEQ ID NO:Y, a polypeptide encoded by the

polynucleotide sequence contained in SEQ ID NO:X or the complement thereof, a polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, a polypeptide encoded by the portion of SEQ ID NO:B as defined in column 6 of Table 1B, and/or a polypeptide encoded by the cDNA contained in Clone ID NO:Z). In particular, N-terminal deletions may be described by the general formula m-q, where q is a whole integer representing the total number of amino acid residues in a polypeptide of the invention (e.g., the polypeptide disclosed in SEQ ID NO:Y, or the polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2), and m is defined as any integer ranging from 2 to q-6. Polynucleotides encoding these polypeptides are also encompassed by the invention.

- residues from the carboxy terminus of the amino acid sequence of a polypeptide disclosed herein (e.g., a polypeptide of SEQ ID NO:Y, a polypeptide encoded by the polynucleotide sequence contained in SEQ ID NO:X, a polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, and/or a polypeptide encoded by the cDNA contained in Clone ID NO:Z). In particular, C-terminal deletions may be described by the general formula 1-n, where n is any whole integer ranging from 6 to q-1, and where n corresponds to the position of amino acid residue in a polypeptide of the invention. Polynucleotides encoding these polypeptides are also encompassed by the invention.
- In addition, any of the above described N- or C-terminal deletions can be combined to produce a N- and C-terminal deleted polypeptide. The invention also provides polypeptides having one or more amino acids deleted from both the amino and the carboxyl termini, which may be described generally as having residues m-n of a polypeptide encoded by SEQ ID NO:X (e.g., including, but not limited to, the preferred polypeptide disclosed as SEQ ID NO:Y and the polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2), the cDNA contained in Clone ID NO:Z, and/or the complement thereof, where n and m are integers as described above. Polynucleotides encoding these polypeptides are also encompassed by the invention.
- [0145] Also as mentioned above, even if deletion of one or more amino acids from the C-terminus of a protein results in modification of loss of one or more biological functions of the protein, other functional activities (e.g., biological activities, ability to multimerize, ability to bind a ligand) may still be retained. For example the ability of the shortened

mutein to induce and/or bind to antibodies which recognize the complete or mature forms of the polypeptide generally will be retained when less than the majority of the residues of the complete or mature polypeptide are removed from the C-terminus. Whether a particular polypeptide lacking C-terminal residues of a complete polypeptide retains such immunologic activities can readily be determined by routine methods described herein and otherwise known in the art. It is not unlikely that a mutein with a large number of deleted C-terminal amino acid residues may retain some biological or immunogenic activities. In fact, peptides composed of as few as six amino acid residues may often evoke an immune response.

- [0146] The present application is also directed to proteins containing polypeptides at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to a polypeptide sequence set forth herein. In preferred embodiments, the application is directed to proteins containing polypeptides at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to polypeptides having the amino acid sequence of the specific N- and C-terminal deletions. Polynucleotides encoding these polypeptides are also encompassed by the invention.
- Any polypeptide sequence encoded by, for example, the polynucleotide sequences set forth as SEQ ID NO:X or the complement thereof, (presented, for example, in Tables 1A and 2), the cDNA contained in Clone ID NO:Z, or the polynucleotide sequence as defined in column 6 of Table 1B, may be analyzed to determine certain preferred regions of the polypeptide. For example, the amino acid sequence of a polypeptide encoded by a polynucleotide sequence of SEQ ID NO:X (e.g., the polypeptide of SEQ ID NO:Y and the polypeptide encoded by the portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2) or the cDNA contained in Clone ID NO:Z may be analyzed using the default parameters of the DNASTAR computer algorithm (DNASTAR, Inc., 1228 S. Park St., Madison, WI 53715 USA; http://www.dnastar.com/).
- [0148] Polypeptide regions that may be routinely obtained using the DNASTAR computer algorithm include, but are not limited to, Garnier-Robson alpha-regions, beta-regions, turn-regions, and coil-regions; Chou-Fasman alpha-regions, beta-regions, and turn-regions; Kyte-Doolittle hydrophilic regions and hydrophobic regions; Eisenberg alpha- and beta-amphipathic regions; Karplus-Schulz flexible regions; Emini surface-forming regions; and Jameson-Wolf regions of high antigenic index. Among

highly preferred polynucleotides of the invention in this regard are those that encode polypeptides comprising regions that combine several structural features, such as several (e.g., 1, 2, 3 or 4) of the features set out above.

- [0149] Additionally, Kyte-Doolittle hydrophilic regions and hydrophobic regions, Emini surface-forming regions, and Jameson-Wolf regions of high antigenic index (i.e., containing four or more contiguous amino acids having an antigenic index of greater than or equal to 1.5, as identified using the default parameters of the Jameson-Wolf program) can routinely be used to determine polypeptide regions that exhibit a high degree of potential for antigenicity. Regions of high antigenicity are determined from data by DNASTAR analysis by choosing values which represent regions of the polypeptide which are likely to be exposed on the surface of the polypeptide in an environment in which antigen recognition may occur in the process of initiation of an immune response.
- [0150] Preferred polypeptide fragments of the invention are fragments comprising, or alternatively, consisting of, an amino acid sequence that displays a functional activity (e.g. biological activity) of the polypeptide sequence of which the amino acid sequence is a fragment. By a polypeptide displaying a "functional activity" is meant a polypeptide capable of one or more known functional activities associated with a full-length protein, such as, for example, biological activity, antigenicity, immunogenicity, and/or multimerization, as described herein.
- [0151] Other preferred polypeptide fragments are biologically active fragments. Biologically active fragments are those exhibiting activity similar, but not necessarily identical, to an activity of the polypeptide of the present invention. The biological activity of the fragments may include an improved desired activity, or a decreased undesirable activity.
- [0152] In preferred embodiments, polypeptides of the invention comprise, or alternatively consist of, one, two, three, four, five or more of the antigenic fragments of the polypeptide of SEQ ID NO:Y, or portions thereof. Polynucleotides encoding these polypeptides are also encompassed by the invention.
- [0153] The present invention encompasses polypeptides comprising, or alternatively consisting of, an epitope of: the polypeptide sequence shown in SEQ ID NO:Y; a polypeptide sequence encoded by SEQ ID NO:X or the complementary strand thereto; the polypeptide sequence encoded by the portion of SEQ ID NO:X as defined in columns 8

and 9 of Table 2; the polypeptide sequence encoded by the portion of SEQ ID NO:B as defined in column 6 of Table 1B or the complement thereto; the polypeptide sequence encoded by the cDNA contained in Clone ID NO:Z; or the polypeptide sequence encoded by a polynucleotide that hybridizes to the sequence of SEQ ID NO:X, the complement of the sequence of SEQ ID NO:X, the complement of a portion of SEQ ID NO:X as defined in columns 8 and 9 of Table 2, or the cDNA sequence contained in Clone ID NO:Z under stringent hybridization conditions or alternatively, under lower stringency hybridization as defined *supra*. The present invention further encompasses polynucleotide sequences encoding an epitope of a polypeptide sequence of the invention (such as, for example, the sequence disclosed in SEQ ID NO:X, or a fragment thereof), polynucleotide sequences of the complementary strand of a polynucleotide sequence encoding an epitope of the invention, and polynucleotide sequences which hybridize to the complementary strand under stringent hybridization conditions or alternatively, under lower stringency hybridization conditions defined *supra*.

- In a term "epitopes," as used herein, refers to portions of a polypeptide having antigenic or immunogenic activity in an animal, preferably a mammal, and most preferably in a human. In a preferred embodiment, the present invention encompasses a polypeptide comprising an epitope, as well as the polynucleotide encoding this polypeptide. An "immunogenic epitope," as used herein, is defined as a portion of a protein that elicits an antibody response in an animal, as determined by any method known in the art, for example, by the methods for generating antibodies described *infra*. (See, for example, Geysen et al., Proc. Natl. Acad. Sci. USA 81:3998-4002 (1983)). The term "antigenic epitope," as used herein, is defined as a portion of a protein to which an antibody can immunospecifically bind its antigen as determined by any method well known in the art, for example, by the immunoassays described herein. Immunospecific binding excludes non-specific binding but does not necessarily exclude cross- reactivity with other antigens. Antigenic epitopes need not necessarily be immunogenic.
- [0155] Fragments which function as epitopes may be produced by any conventional means. (See, e.g., Houghten, R. A., Proc. Natl. Acad. Sci. USA 82:5131-5135 (1985) further described in U.S. Patent No. 4,631,211.)
- [0156] In the present invention, antigenic epitopes preferably contain a sequence of at least 4, at least 5, at least 6, at least 7, more preferably at least 8, at least 9, at least 10, at

least 11, at least 12, at least 13, at least 14, at least 15, at least 20, at least 25, at least 30, at least 40, at least 50, and, most preferably, between about 15 to about 30 amino acids. Preferred polypeptides comprising immunogenic or antigenic epitopes are at least 10, 15, 20, 25, 30, 35, 40, 45, 50, 55, 60, 65, 70, 75, 80, 85, 90, 95, or 100 amino acid residues in length. Additional non-exclusive preferred antigenic epitopes include the antigenic epitopes disclosed herein, as well as portions thereof. Antigenic epitopes are useful, for example, to raise antibodies, including monoclonal antibodies, that specifically bind the epitope. Preferred antigenic epitopes include the antigenic epitopes disclosed herein, as well as any combination of two, three, four, five or more of these antigenic epitopes. Antigenic epitopes can be used as the target molecules in immunoassays. (See, for instance, Wilson et al., Cell 37:767-778 (1984); Sutcliffe et al., Science 219:660-666 (1983)).

Non-limiting examples of epitopes of polypeptides that can be used to generate [0157] antibodies of the invention include a polypeptide comprising, or alternatively consisting of, at least one, two, three, four, five, six or more of the portion(s) of SEQ ID NO:Y specified in column 6 of Table 1A. These polypeptide fragments have been determined to bear antigenic epitopes of the proteins of the invention by the analysis of the Jameson-Wolf antigenic index which is included in the DNAStar suite of computer programs. By "comprise" it is intended that a polypeptide contains at least one, two, three, four, five, six or more of the portion(s) of SEQ ID NO:Y shown in column 6 of Table 1A, but it may contain additional flanking residues on either the amino or carboxyl termini of the recited portion. Such additional flanking sequences are preferably sequences naturally found adjacent to the portion; i.e., contiguous sequence shown in SEQ ID NO:Y. The flanking sequence may, however, be sequences from a heterologous polypeptide, such as from another protein described herein or from a heterologous polypeptide not described herein. In particular embodiments, epitope portions of a polypeptide of the invention comprise one, two, three, or more of the portions of SEQ ID NO:Y shown in column 6 of Table 1A. Polynucleotides encoding these polypeptides are also encompassed by the invention.

[0158] Similarly, immunogenic epitopes can be used, for example, to induce antibodies according to methods well known in the art. See, for instance, Sutcliffe et al., supra; Wilson et al., supra; Chow et al., Proc. Natl. Acad. Sci. USA 82:910-914; and Bittle et al.; J. Gen. Virol. 66:2347-2354 (1985). Preferred immunogenic epitopes

include the immunogenic epitopes disclosed herein, as well as any combination of two, three, four, five or more of these immunogenic epitopes. The polypeptides comprising one or more immunogenic epitopes may be presented for eliciting an antibody response together with a carrier protein, such as an albumin, to an animal system (such as rabbit or mouse), or, if the polypeptide is of sufficient length (at least about 25 amino acids), the polypeptide may be presented without a carrier. However, immunogenic epitopes comprising as few as 8 to 10 amino acids have been shown to be sufficient to raise antibodies capable of binding to, at the very least, linear epitopes in a denatured polypeptide (e.g., in Western blotting).

[0159] Epitope-bearing polypeptides of the present invention may be used to induce antibodies according to methods well known in the art including, but not limited to, in vivo immunization, in vitro immunization, and phage display methods. See, e.g., Sutcliffe et al., supra; Wilson et al., supra, and Bittle et al., J. Gen. Virol., 66:2347-2354 (1985). If in vivo immunization is used, animals may be immunized with free peptide; however, antipeptide antibody titer may be boosted by coupling the peptide to a macromolecular carrier, such as keyhole limpet hemacyanin (KLH) or tetanus toxoid. For instance, peptides containing cysteine residues may be coupled to a carrier using a linker such as maleimidobenzoyl- N-hydroxysuccinimide ester (MBS), while other peptides may be coupled to carriers using a more general linking agent such as glutaraldehyde. Animals such as rabbits, rats and mice are immunized with either free or carrier- coupled peptides, for instance, by intraperitoneal and/or intradermal injection of emulsions containing about 100 µg of peptide or carrier protein and Freund's adjuvant or any other adjuvant known for stimulating an immune response. Several booster injections may be needed, for instance, at intervals of about two weeks, to provide a useful titer of anti-peptide antibody which can be detected, for example, by ELISA assay using free peptide adsorbed to a solid surface. The titer of anti-peptide antibodies in serum from an immunized animal may be increased by selection of anti-peptide antibodies, for instance, by adsorption to the peptide on a solid support and elution of the selected antibodies according to methods well known in the art.

[0160] As one of skill in the art will appreciate, and as discussed above, the polypeptides of the present invention (e.g., those comprising an immunogenic or antigenic epitope) can be fused to heterologous polypeptide sequences. For example, polypeptides of the present

invention (including fragments or variants thereof), may be fused with the constant domain of immunoglobulins (IgA, IgE, IgG, IgM), or portions thereof (CH1, CH2, CH3, or any combination thereof and portions thereof, resulting in chimeric polypeptides. By way of another non-limiting example, polypeptides and/or antibodies of the present invention (including fragments or variants thereof) may be fused with albumin (including but not limited to recombinant human serum albumin or fragments or variants thereof (see, e.g., U.S. Patent No. 5,876,969, issued March 2, 1999, EP Patent 0 413 622, and U.S. Patent No. 5,766,883, issued June 16, 1998, herein incorporated by reference in their entirety)). In a preferred embodiment, polypeptides and/or antibodies of the present invention (including fragments or variants thereof) are fused with the mature form of human serum albumin (i.e., amino acids 1 - 585 of human serum albumin as shown in Figures 1 and 2 of EP Patent 0 322 094) which is herein incorporated by reference in its entirety. In another preferred embodiment, polypeptides and/or antibodies of the present invention (including fragments or variants thereof) are fused with polypeptide fragments comprising, or alternatively consisting of, amino acid residues 1-z of human serum albumin, where z is an integer from 369 to 419, as described in U.S. Patent 5,766,883 herein incorporated by reference in its entirety. Polypeptides and/or antibodies of the present invention (including fragments or variants thereof) may be fused to either the Nor C-terminal end of the heterologous protein (e.g., immunoglobulin Fc polypeptide or human serum albumin polypeptide). Polynucleotides encoding fusion proteins of the invention are also encompassed by the invention.

[0161] Such fusion proteins as those described above may facilitate purification and may increase half-life *in vivo*. This has been shown for chimeric proteins consisting of the first two domains of the human CD4-polypeptide and various domains of the constant regions of the heavy or light chains of mammalian immunoglobulins. See, e.g., EP 394,827; Traunecker et al., Nature, 331:84-86 (1988). Enhanced delivery of an antigen across the epithelial barrier to the immune system has been demonstrated for antigens (e.g., insulin) conjugated to an FcRn binding partner such as IgG or Fc fragments (see, e.g., PCT Publications WO 96/22024 and WO 99/04813). IgG Fusion proteins that have a disulfide-linked dimeric structure due to the IgG portion desulfide bonds have also been found to be more efficient in binding and neutralizing other molecules than monomeric polypeptides or fragments thereof alone. See, e.g., Fountoulakis et al., J. Biochem.,

270:3958-3964 (1995). Nucleic acids encoding the above epitopes can also be recombined with a gene of interest as an epitope tag (e.g., the hemagglutinin (HA) tag or flag tag) to aid in detection and purification of the expressed polypeptide. For example, a system described by Janknecht et al. allows for the ready purification of non-denatured fusion proteins expressed in human cell lines (Janknecht et al., 1991, Proc. Natl. Acad. Sci. USA 88:8972-897). In this system, the gene of interest is subcloned into a vaccinia recombination plasmid such that the open reading frame of the gene is translationally fused to an amino-terminal tag consisting of six histidine residues. The tag serves as a matrix binding domain for the fusion protein. Extracts from cells infected with the recombinant vaccinia virus are loaded onto Ni2+ nitriloacetic acid-agarose column and histidine-tagged proteins can be selectively eluted with imidazole-containing buffers.

Fusion Proteins

- [0162] Any polypeptide of the present invention can be used to generate fusion proteins. For example, the polypeptide of the present invention, when fused to a second protein, can be used as an antigenic tag. Antibodies raised against the polypeptide of the present invention can be used to indirectly detect the second protein by binding to the polypeptide. Moreover, because secreted proteins target cellular locations based on trafficking signals, polypeptides of the present invention which are shown to be secreted can be used as targeting molecules once fused to other proteins.
- [0163] Examples of domains that can be fused to polypeptides of the present invention include not only heterologous signal sequences, but also other heterologous functional regions. The fusion does not necessarily need to be direct, but may occur through linker sequences.
- [0164] In certain preferred embodiments, proteins of the invention are fusion proteins comprising an amino acid sequence that is an N and/or C- terminal deletion of a polypeptide of the invention. In preferred embodiments, the invention is directed to a fusion protein comprising an amino acid sequence that is at least 80%, 85%, 90%, 95%, 96%, 97%, 98% or 99% identical to a polypeptide sequence of the invention. Polynucleotides encoding these proteins are also encompassed by the invention.
- [0165] Moreover, fusion proteins may also be engineered to improve characteristics of the polypeptide of the present invention. For instance, a region of additional amino acids,

particularly charged amino acids, may be added to the N-terminus of the polypeptide to improve stability and persistence during purification from the host cell or subsequent handling and storage. Also, peptide moieties may be added to the polypeptide to facilitate purification. Such regions may be removed prior to final preparation of the polypeptide. The addition of peptide moieties to facilitate handling of polypeptides are familiar and routine techniques in the art.

[0166] As one of skill in the art will appreciate that, as discussed above, polypeptides of the present invention, and epitope-bearing fragments thereof, can be combined with heterologous polypeptide sequences. For example, the polypeptides of the present invention may be fused with heterologous polypeptide sequences, for example, the polypeptides of the present invention may be fused with the constant domain of immunoglobulins (IgA, IgE, IgG, IgM) or portions thereof (CH1, CH2, CH3, and any combination thereof, including both entire domains and portions thereof), or albumin (including, but not limited to, native or recombinant human albumin or fragments or variants thereof (see, e.g., U.S. Patent No. 5,876,969, issued March 2, 1999, EP Patent 0 413 622, and U.S. Patent No. 5,766,883, issued June 16, 1998, herein incorporated by reference in their entirety)), resulting in chimeric polypeptides. For example, EP-A-O 464 533 (Canadian counterpart 2045869) discloses fusion proteins comprising various portions of constant region of immunoglobulin molecules together with another human protein or part thereof. In many cases, the Fc part in a fusion protein is beneficial in therapy and diagnosis, and thus can result in, for example, improved pharmacokinetic properties (EP-A 0232 262). Alternatively, deleting the Fc part after the fusion protein has been expressed, detected, and purified, would be desired. For example, the Fc portion may hinder therapy and diagnosis if the fusion protein is used as an antigen for immunizations. In drug discovery, for example, human proteins, such as hIL-5, have been fused with Fc portions for the purpose of high-throughput screening assays to identify antagonists of hIL-5. See, D. Bennett et al., J. Molecular Recognition 8:52-58 (1995); K. Johanson et al., J. Biol. Chem. 270:9459-9471 (1995).

[0167] Moreover, the polypeptides of the present invention can be fused to marker sequences, such as a polypeptide which facilitates purification of the fused polypeptide. In preferred embodiments, the marker amino acid sequence is a hexa-histidine peptide, such as the tag provided in a pQE vector (QIAGEN, Inc., 9259 Eton Avenue, Chatsworth,

CA, 91311), among others, many of which are commercially available. As described in Gentz et al., Proc. Natl. Acad. Sci. USA 86:821-824 (1989), for instance, hexa-histidine provides for convenient purification of the fusion protein. Another peptide tag useful for purification, the "HA" tag, corresponds to an epitope derived from the influenza hemagglutinin protein (Wilson et al., Cell 37:767 (1984).)

[0168] Additional fusion proteins of the invention may be generated through the techniques of gene-shuffling, motif-shuffling, exon-shuffling, and/or codon-shuffling (collectively referred to as "DNA shuffling"), briefly described below, and further described herein. DNA shuffling may be employed to modulate the activities of polypeptides of the invention, such methods can be used to generate polypeptides with altered activity, as well as agonists and antagonists of the polypeptides. See, generally, U.S. Patent Nos. 5,605,793; 5,811,238; 5,830,721; 5,834,252; and 5,837,458, and Patten et al., Curr. Opinion Biotechnol. 8:724-33 (1997); Harayama, Trends Biotechnol. 16(2):76-82 (1998); Hansson et al., J. Mol. Biol. 287:265-76 (1999); and Lorenzo and Blasco, Biotechniques 24(2):308-13 (1998); each of these patents and publications are hereby incorporated by reference in its entirety). In a preferred embodiment, one or more components, motifs, sections, parts, domains, fragments, etc., of a polynucleotide encoding a polypeptide of the invention may be recombined with one or more components, motifs, sections, parts, domains, fragments, etc., of one or more heterologous molecules encoding a heterologous polypeptide.

[0169] Thus, any of these above fusions can be engineered using the polynucleotides or the polypeptides of the present invention.

Recombinant and Synthetic Production of Polypeptides of the Invention

- [0170] The present invention also relates to vectors containing the polynucleotide of the present invention, host cells, and the production of polypeptides by synthetic and recombinant techniques. The vector may be, for example, a phage, plasmid, viral, or retroviral vector. Retroviral vectors may be replication competent or replication defective. In the latter case, viral propagation generally will occur only in complementing host cells.
- [0171] The polynucleotides of the invention may be joined to a vector containing a selectable marker for propagation in a host. Generally, a plasmid vector is introduced in a precipitate, such as a calcium phosphate precipitate, or in a complex with a charged lipid.

If the vector is a virus, it may be packaged in vitro using an appropriate packaging cell line and then transduced into host cells.

[0172] The polynucleotide insert should be operatively linked to an appropriate promoter, such as the phage lambda PL promoter, the E. coli lac, trp, phoA and tac promoters, the SV40 early and late promoters and promoters of retroviral LTRs, to name a few. Other suitable promoters will be known to the skilled artisan. The expression constructs will further contain sites for transcription initiation, termination, and, in the transcribed region, a ribosome binding site for translation. The coding portion of the transcripts expressed by the constructs will preferably include a translation initiating codon at the beginning and a termination codon (UAA, UGA or UAG) appropriately positioned at the end of the polypeptide to be translated.

[0173] As indicated, the expression vectors will preferably include at least one selectable marker. Such markers include dihydrofolate reductase, G418 or neomycin resistance, glutamine synthase, for eukaryotic cell culture and tetracycline, kanamycin or ampicillin resistance genes for culturing in *E. coli* and other bacteria. Representative examples of appropriate hosts include, but are not limited to, bacterial cells, such as *E. coli*, Streptomyces and Salmonella typhimurium cells; fungal cells, such as yeast cells (e.g., Saccharomyces cerevisiae or Pichia pastoris (ATCC Accession No. 201178)); insect cells such as Drosophila S2 and Spodoptera Sf9 cells; animal cells such as CHO, COS, 293, NSO and Bowes melanoma cells; and plant cells. Appropriate culture mediums and conditions for the above-described host cells are known in the art.

[0174] Among vectors preferred for use in bacteria include pQE70, pQE60 and pQE-9, available from QIAGEN, Inc.; pBluescript vectors, Phagescript vectors, pNH8A, pNH16a, pNH18A, pNH46A, available from Stratagene Cloning Systems, Inc.; and ptrc99a, pKK223-3, pKK233-3, pDR540, pRIT5 available from Pharmacia Biotech, Inc. Among preferred eukaryotic vectors are pWLNEO, pSV2CAT, pOG44, pXT1 and pSG available from Stratagene; and pSVK3, pBPV, pMSG and pSVL available from Pharmacia. Preferred expression vectors for use in yeast systems include, but are not limited to pYES2, pYD1, pTEF1/Zeo, pYES2/GS, pPICZ, pGAPZ, pGAPZalph, pPIC9, pPIC3.5, pHIL-D2, pHIL-S1, pPIC3.5K, pPIC9K, and PAO815 (all available from Invitrogen, Carlsbad, CA). Other suitable vectors will be readily apparent to the skilled artisan.

Vectors which use glutamine synthase (GS) or DHFR as the selectable markers [0175]can be amplified in the presence of the drugs methionine sulphoximine or methotrexate, respectively. An advantage of glutamine synthase based vectors are the availabilty of cell lines (e.g., the murine myeloma cell line, NS0) which are glutamine synthase negative. Glutamine synthase expression systems can also function in glutamine synthase expressing cells (e.g., Chinese Hamster Ovary (CHO) cells) by providing additional inhibitor to prevent the functioning of the endogenous gene. A glutamine synthase expression system and components thereof are detailed in PCT publications: WO87/04462; WO86/05807; WO89/01036; WO89/10404; and WO91/06657 which are hereby incorporated in their entireties by reference herein. Additionally, glutamine synthase expression vectors can be obtained from Lonza Biologics, Inc. (Portsmouth, NH). Expression and production of monoclonal antibodies using a GS expression system in murine myeloma cells is described in Bebbington et al., Bio/technology 10:169(1992) and in Biblia and Robinson Biotechnol. Prog. 11:1 (1995) which are herein incorporated by reference.

- [0176] The present invention also relates to host cells containing the above-described vector constructs described herein, and additionally encompasses host cells containing nucleotide sequences of the invention that are operably associated with one or more heterologous control regions (e.g., promoter and/or enhancer) using techniques known of in the art. The host cell can be a higher eukaryotic cell, such as a mammalian cell (e.g., a human derived cell), or a lower eukaryotic cell, such as a yeast cell, or the host cell can be a prokaryotic cell, such as a bacterial cell. A host strain may be chosen which modulates the expression of the inserted gene sequences, or modifies and processes the gene product in the specific fashion desired. Expression from certain promoters can be elevated in the presence of certain inducers; thus expression of the genetically engineered polypeptide may be controlled. Furthermore, different host cells have characteristics and specific mechanisms for the translational and post-translational processing and modification (e.g., phosphorylation, cleavage) of proteins. Appropriate cell lines can be chosen to ensure the desired modifications and processing of the foreign protein expressed.
- [0177] Introduction of the nucleic acids and nucleic acid constructs of the invention into the host cell can be effected by calcium phosphate transfection, DEAE-dextran mediated transfection, cationic lipid-mediated transfection, electroporation, transduction,

infection, or other methods. Such methods are described in many standard laboratory manuals, such as Davis et al., Basic Methods In Molecular Biology (1986). It is specifically contemplated that the polypeptides of the present invention may in fact be expressed by a host cell lacking a recombinant vector.

[0178] In addition to encompassing host cells containing the vector constructs discussed herein, the invention also encompasses primary, secondary, and immortalized host cells of vertebrate origin, particularly mammalian origin, that have been engineered to delete or replace endogenous genetic material (e.g., immune/hematopoietic antigen coding sequence), and/or to include genetic material (e.g., heterologous polynucleotide sequences) that is operably associated with immune/hematopoietic associated polynucleotides of the invention, and which activates, alters, and/or amplifies endogenous immune/hematopoietic associated polynucleotides. For example, techniques known in the art may be used to operably associate heterologous control regions (e.g., promoter and/or enhancer) and endogenous immune/hematopoietic associated polynucleotide sequences via homologous recombination (see, e.g., U.S. Patent Number 5,641,670, issued June 24, 1997; International Publication Number WO 96/29411; International Publication Number WO 94/12650; Koller et al., Proc. Natl. Acad. Sci. USA 86:8932-8935 (1989); and Zijlstra et al., Nature 342:435-438 (1989), the disclosures of each of which are incorporated by reference in their entireties).

Polypeptides of the present invention can also be recovered from: products purified from natural sources, including bodily fluids, tissues and cells, whether directly isolated or cultured; products of chemical synthetic procedures; and products produced by recombinant techniques from a prokaryotic or eukaryotic host, including, for example, bacterial, yeast, higher plant, insect, and mammalian cells. Depending upon the host employed in a recombinant production procedure, the polypeptides of the present invention may be glycosylated or may be non-glycosylated. In addition, polypeptides of the invention may also include an initial modified methionine residue, in some cases as a result of host-mediated processes. Thus, it is well known in the art that the N-terminal methionine encoded by the translation initiation codon generally is removed with high efficiency from any protein after translation in all eukaryotic cells. While the N-terminal methionine on most proteins also is efficiently removed in most prokaryotes, for some

proteins, this prokaryotic removal process is inefficient, depending on the nature of the amino acid to which the N-terminal methionine is covalently linked.

- [0180] In one embodiment, the yeast Pichia pastoris is used to express polypeptides of the invention in a eukaryotic system. Pichia pastoris is a methylotrophic yeast which can metabolize methanol as its sole carbon source. A main step in the methanol metabolization pathway is the oxidation of methanol to formaldehyde using O₂. This reaction is catalyzed by the enzyme alcohol oxidase. In order to metabolize methanol as its sole carbon source, Pichia pastoris must generate high levels of alcohol oxidase due, in part, to the relatively low affinity of alcohol oxidase for O₂. Consequently, in a growth medium depending on methanol as a main carbon source, the promoter region of one of the two alcohol oxidase genes (AOXI) is highly active. In the presence of methanol, alcohol oxidase produced from the AOXI gene comprises up to approximately 30% of the total soluble protein in Pichia pastoris. See, Ellis, S.B., et al., Mol. Cell. Biol. 5:1111-21 (1985); Koutz, P.J, et al., Yeast 5:167-77 (1989); Tschopp, J.F., et al., Nucl. Acids Res. 15:3859-76 (1987). Thus, a heterologous coding sequence, such as, for example, a polynucleotide of the present invention, under the transcriptional regulation of all or part of the AOXI regulatory sequence is expressed at exceptionally high levels in Pichia yeast grown in the presence of methanol.
- In one example, the plasmid vector pPIC9K is used to express DNA encoding a polypeptide of the invention, as set forth herein, in a *Pichea* yeast system essentially as described in "*Pichia* Protocols: Methods in Molecular Biology," D.R. Higgins and J. Cregg, eds. The Humana Press, Totowa, NJ, 1998. This expression vector allows expression and secretion of a polypeptide of the invention by virtue of the strong *AOX1* promoter linked to the *Pichia pastoris* alkaline phosphatase (PHO) secretory signal peptide (i.e., leader) located upstream of a multiple cloning site.
- [0182] Many other yeast vectors could be used in place of pPIC9K, such as, pYES2, pYD1, pTEF1/Zeo, pYES2/GS, pPICZ, pGAPZ, pGAPZalpha, pPIC9, pPIC3.5, pHIL-D2, pHIL-S1, pPIC3.5K, and PAO815, as one skilled in the art would readily appreciate, as long as the proposed expression construct provides appropriately located signals for